

SEQUENCE LISTING

:110~ Sternberg, Paul W.
Barr, Maureen M.

:120~ POLYCYSTIC KIDNEY DISEASE GENE HOMOLOGS REQUIRED FOR MALE MATING BEHAVIOR IN NEMATODES AND ASSAYS BASED THEREON

:130~ 18021 2901B

:140~ Unassigned

:141~ 2900-09-05

:150~ 09/479,467

:151~ 2900-01-06

:160~ 60-115,127

:151~ 1999-01-06

:160~ 16

:170~ Patent In Ver. 2.0

:210~ 1
:211~ 12912
:212~ DNA
:213~ Homo sapiens PKD-1 gene

:220~
:221~ CDS
:222~ (1)...(12912)

:400~ 1
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Met Pro Pro Ala Ala Pro Ala Arg Leu Ala Leu Ala Leu Gly Leu Gly
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ctg tgg ctc ggg gcg ctg gcg ggg ccc ggg cgc ggc tgc ggg ccc 96
Leu Trp Leu Gly Ala Leu Ala Gly Gly Pro Gly Arg Gly Cys Gly Pro
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tgc gag ccc ccc tgc ctc tgc ggg cca gcg ccc ggc gcc gcc tgc cgc 144
Cys Glu Pro Pro Cys Leu Cys Gly Pro Ala Pro Gly Ala Ala Cys Arg
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gtc aac tgc tcg ggc cgc ggg ctg cgg acg ctc ggt ccc ggg ctg cgc 192
Val Asn Cys Ser Gly Arg Gly Leu Arg Thr Leu Gly Pro Ala Leu Arg
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atc ccc gcg gac gcc aca gag cta gac gtc tcc cac aac ctg ctc cgg 240
Ile Pro Ala Asp Ala Thr Glu Leu Asp Val Ser His Asn Leu Leu Arg
65 70 75 80

gcg ctg gac gtt ggg ctc ctg gcg aac ctc tgc gcg ctg gca gag ctg 288
Ala Leu Asp Val Gly Leu Leu Ala Asn Leu Ser Ala Leu Ala Glu Leu
85 90 95

gat ata agc aac aac aag att tct acg tta gaa gaa gga ata ttt gct 336
Asp Ile Ser Asn Asn Lys Ile Ser Thr Leu Glu Glu Gly Ile Phe Ala
100 105 110

aat tta ttt aat tta agt gaa ata aac ctg agt ggg aac ccg ttt gag 384
Asn Leu Phe Asn Leu Ser Glu Ile Asn Leu Ser Gly Asn Pro Phe Glu
115 120 125

tgt gac tgt ggc ctg gcg tgg ctg ccg caa tgg gcg gag gag cag cag 432
Cys Asp Cys Gly Leu Ala Trp Leu Pro Gln Trp Ala Glu Glu Gln Gln

130	135	140	
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ggc gag gag tat gtc gca tcc cct gag aac aac tca ggc acc gtc Gly Glu Tyr Val Ala Cys Leu Pro Asp Asn Ser Ser Gly Thr Val 180 185 190			576
gca gca gtc tcc ttt tca gct gcc cas gaa ggc ctg ctt cag cca gag Ala Ala Val Ser Phe Ser Ala Ala His Glu Gly Leu Leu Gln Pro Glu 195 200 205			624
gcc tgc aac gcc ttc tgc ttc tcc acc ggc cag ggc ctc gca gcc ctc Ala Cys Ser Ala Phe Cys Phe Ser Thr Gly Gln Gly Leu Ala Ala Leu 210 215 220			671
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ccc acc tgt agg ggc ccc acc ctc ctc cag cac gtc ttc cct gcc tcc Pro Thr Cys Arg Gly Pro Thr Leu Leu Gln His Val Phe Pro Ala Ser 260 265 270			816
cca ggg gcc acc ctg gtg ggg ccc cas gga cct ctg gcc tct ggc cag Pro Gly Ala Thr Leu Val Gly Pro His Gly Pro Leu Ala Ser Gly Gln 275 280 285			864
cta gca gcc ttc cac atc gct gcc ccc ctc cct gtc act gac aca cgc Leu Ala Ala Phe His Ile Ala Ala Pro Leu Pro Val Thr Asp Thr Arg 290 295 300			912
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gcc tcg cat cgc tat gtg ctg cct ggg cgc tat cac gtg acg gcc gtg Ala Ser His Arg Tyr Val Leu Pro Gly Arg Tyr His Val Thr Ala Val 325 330 335			1008
ctg gcc ctg ggg gcc ggc tca gcc ctg ctg ggg aca gac gtg cag gtg Leu Ala Leu Gly Ala Gly Ser Ala Leu Leu Gly Thr Asp Val Gln Val 340 345 350			1056
gaa gcg gca cct gcc gcc ctg gag ctc gtg tgc ccg tcc tcg gtg cag Glu Ala Ala Pro Ala Ala Leu Glu Leu Val Cys Pro Ser Ser Val Gln 355 360 365			1104
agt gac gag agc ctc gac ctc acg atc cag aac cgc ggt ggt tca ggc Ser Asp Glu Ser Leu Asp Leu Ser Ile Gln Asn Arg Gly Gly Ser Gly 370 375 380			1152
ctg gag gcc gcc tac agc atc gtg gcc ctg ggc gag gag ccg gcc cga Leu Glu Ala Ala Tyr Ser Ile Val Ala Leu Gly Glu Glu Pro Ala Arg 385 390 395 400			1200
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Ala Val His Pro Leu Cys Pro Ser Asp Thr Glu Ile Phe Pro Gly Asn
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 Gly His Cys Tyr Arg Leu Val Val Glu Lys Ala Ala Trp Leu Gln Ala
 420 425 430
 cag gag caa tgt cag gcc tgg ggg gcc gcc ctg gca atg gtg gac 1344
 Gln Glu Gln Cys Gln Ala Trp Ala Gly Ala Ala Leu Ala Met Val Asp
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 agt ccc gcc gtg cag cgc ttc ctg gtc tcc cgg gtc acc agg aac cta 1392
 Ser Pro Ala Val Gln Arg Phe Leu Val Ser Arg Val Thr Arg Ser Leu
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 gag gtg tgg atc ggc ttc tcg act gtg cag ggg gtg gag gtg ggc cca 1440
 Asp Val Trp Ile Gly Phe Ser Thr Val Gln Gly Val Glu Val Gly Pro
 465 470 475 480
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 Ala Pro Gln Gly Glu Ala Phe Ser Leu Glu Ser Cys Gln Asn Trp Leu
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 Pro Gly Glu Pro His Pro Ala Thr Ala Glu His Cys Val Arg Leu Gly
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 Pro Thr Gly Trp Cys Asn Thr Asp Leu Cys Ser Ala Pro His Ser Tyr
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 Leu Val Gly Ala Pro Ser Gly Asp Leu Gln Gly Pro Leu Thr Pro Leu
 545 550 555 560
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 Ala Gln Gln Asp Gly Leu Ser Ala Pro His Glu Pro Val Glu Val Met
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 Val Phe Pro Gly Leu Arg Leu Ser Arg Glu Ala Phe Leu Thr Thr Ala
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 Val Tyr Arg Leu Leu Ser Thr Ala Gly Thr Pro Glu Asn Gly Ser Glu
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 Pro Glu Ser Arg Ser Pro Asp Asn Arg Thr Gln Leu Ala Pro Ala Cys
 625 630 635 640
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 645 650 655
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 Asp Ala Ser Cys His Pro Gln Ala Cys Ala Asn Gly Cys Thr Ser Gly
 660 665 670

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Gln	Asp	Val	Ieu	Met	Leu	Pro	Gly	Asp	Leu	Val	Gly	Ieu	Gln	His	Asp		
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Ala	Gly	Pro	Gly	Ala	Ieu	Leu	His	Cys	Ser	Pro	Ala	Pro	Gly	His	Pro		
725								730						735			
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Gly	Pro	Arg	Ala	Pro	Tyr	Ieu	Ser	Ala	Asn	Ala	Ser	Ser	Trp	Ieu	Pro		
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Val	Val	Ser	Pro	Val	Ala	Gly	Ieu	Arg	Val	Ile	Tyr	Pro	Ala	Pro	Arg		
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Asp	Gly	Arg	Ieu	Tyr	Val	Pro	Thr	Asn	Gly	Ser	Ala	Ieu	Val	Ieu	Gln		
835								840						845			
gtg	gac	tct	gg	aa	cc	gg	cc	ac	cc	gt	cc	tt	gg	cc	gg		2592
Val	Asp	Ser	Gly	Ala	Asn	Ala	Thr	Ala	Thr	Ala	Arg	Trp	Pro	Gly	Gly		
850								855						860			
agt	ctc	agc	gcc	ccg	ttt	gag	aat	gtc	tcc	cct	gcc	ctg	gtg	gcc	acc		2640
Ser	Leu	Ser	Ala	Arg	Phe	Glu	Asn	Val	Cys	Pro	Ala	Ieu	Val	Ala	Thr		
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Phe	Val	Pro	Ala	Cys	Pro	Trp	Glu	Thr	Asn	Asp	Thr	Leu	Phe	Ser	Val		
885								890						895			
gta	gca	ctg	ccg	tgg	ctc	agt	gag	ggg	gag	cac	gtg	gtg	gac	gtg	gtg		2736
Val	Ala	Ieu	Pro	Trp	Leu	Ser	Glu	Gly	Glu	His	Val	Val	Asp	Val	Val		
900								905						910			
gtg	gaa	aac	agc	gcc	agc	egg	ggc	aac	ctc	agc	ctg	ccg	gtg	acg	ccg		2784
Val	Glu	Asn	Ser	Ala	Ser	Arg	Ala	Asn	Leu	Ser	Leu	Arg	Val	Thr	Ala		
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930								935						940			

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Phe	Gln	Asn	Val	Val	Phe	Asn	Val	Ile	Tyr	Gln	Ser	Ala	Ala	Val	Phe		
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aag	ttc	tca	ctg	acc	ggc	tcc	aac	cac	gtg	agg	aaa	gtc	acc	gtg	aac		3024
Lys	Leu	Ser	Leu	Ihr	Ala	Ser	Asn	His	Val	Ser	Asn	Val	Ihr	Val	Asn		
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taa	aac	gta	acc	gtg	gag	cgg	atg	aac	agg	atg	cag	gtt	gtg	cag	gtc		3072
Tyr	Asn	Val	Ihr	Val	Glu	Arg	Met	Asn	Arg	Met	Gln	Gly	Leu	Gln	Val		
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Ser	Ihr	Val	Pro	Ala	Val	Ser	Pro	Asn	Ala	Ihr	Leu	Ala	Leu	Thr			
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ccg	ggc	gtg	ctg	gtg	gac	tcc	gcc	gtg	gag	gtg	gcc	tcc	ctg	tgg	acc		3163
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Phe	Gly	Asp	Gly	Glu	Gln	Ala	Leu	His	Gln	Phe	Gln	Pro	Pro	Tyr	Asn		
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gag	tcc	tcc	cca	gtt	cca	gac	ccc	tcc	gtg	gcc	cag	gtg	ctg	gtg	gag		3264
Glu	Ser	Phe	Pro	Val	Pro	Asp	Pro	Ser	Val	Ala	Gln	Val	Leu	Val	Glu		
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cac	aat	gtc	acc	cac	acc	tac	gct	gcc	cca	ggt	gag	tac	ctc	ctg	acc		3312
His	Asn	Val	Thr	His	Thr	Tyr	Ala	Ala	Pro	Gly	Glu	Tyr	Leu	Leu	Thr		
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gtg	ctg	gca	tct	aat	gcc	tcc	gag	aac	ctg	acc	cag	cag	gtg	cct	gtg		3360
Val	Leu	Ala	Ser	Asn	Ala	Phe	Glu	Asn	Leu	Thr	Gln	Gln	Val	Pro	Val		
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agc	gtg	ccg	gcc	tcc	ctg	ccc	tcc	gtg	gct	gtg	ggt	gtg	agt	gac	ggc		3408
Ser	Val	Arg	Ala	Ser	Leu	Pro	Ser	Val	Ala	Val	Gly	Val	Ser	Asp	Gly		
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gtc	ctg	gtg	gcc	ggc	ccg	ccc	gtc	acc	tcc	tac	ccg	cac	ccg	ctg	ccc		3456
Val	Leu	Val	Ala	Gly	Arg	Pro	Val	Thr	Phe	Tyr	Pro	His	Pro	Leu	Pro		
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tcg	cct	ggg	ggt	gtt	ttt	tac	acg	tgg	gac	tcc	ggg	gac	ggc	tcc	cct		3504
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Val	Leu	Thr	Gln	Ser	Gln	Pro	Ala	Ala	Asn	His	Thr	Tyr	Ala	Ser	Arg		
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ggc	acc	tac	cac	gtc	ccg	ctg	gag	gtc	aac	aac	acg	gtg	agg	ggt	ggc		3600
Gly	Thr	Tyr	His	Val	Arg	Leu	Glu	Val	Asn	Asn	Thr	Val	Ser	Gly	Ala		
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gcg	gcc	cag	cac	gtc	ccg	gtc	ttt	gag	gag	ctc	ccg	gga	ctc	agc		3648	
Ala	Ala	Gln	Ala	Asp	Val	Phe	Glu	Glu	Glu	Leu	Arg	Gly	Leu	Ser			

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ggc gcg gtg caa acg ggc gag aac atc acg tgg acc ttc gag atg ggg Ala Ala Val Gln Thr Gly Asp Asn Ile Thr Trp Thr Phe Asp Met Gly			3744
1235	1240	1245	
gaa ggc aac gtg ctg tgg ggc ccc gag gca aca gtg gag cat gtg tac Asp Gly Thr Val Leu Ser Gly Pro Glu Ala Thr Val Glu His Val Tyr			3792
1250	1255	1260	
ctg cgg gca cag aac tgc aca gtg acc gtg ggt gcg ggc aac ccc gcc Leu Arg Ala Gln Asn Cys Thr Val Thr Val Gly Ala Gly Ser Pro Ala			3840
1265	1270	1275	1280
ggc cac ctg gcc cgg aac ctg cac gtg ctg gtc ttc gtc stg gag gtg Gly His Leu Ala Arg Ser Leu His Val Leu Val Phe Val Leu Glu Val			3888
1285	1290	1295	
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ctc acg gcc tac gtc acc ggg aac ccc gac cac tac ctc ttc gag tgg Leu Thr Ala Tyr Val Thr Gly Asn Pro Ala His Tyr Leu Phe Asp Trp			3984
1315	1320	1325	
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1395	1400	1405	
ccc tac cgc tac acc tgg gac ttt ggc acc gag gaa gcc gcc ccc acc Pro Tyr Arg Tyr Thr Trp Asp Phe Gly Thr Glu Ala Ala Pro Thr			4272
1410	1415	1420	
cgt gcc agg ggc cct gag gtg acg ttc atc tac cga gac cca ggc tcc Arg Ala Arg Gly Pro Glu Val Thr Phe Ile Tyr Arg Asp Pro Gly Ser			4320
1425	1430	1435	1440
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1445	1450	1455	
tca gcc ctg gtg gag gtg cag gag ccc gtg ctg gtc acc acg atc aag Ser Ala Leu Val Glu Val Gln Glu Pro Val Leu Val Thr Ser Ile Lys			4416
1460	1465	1470	
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Val Asn Gly Ser Leu Gly Leu Glu Leu Gln Gln Pro Tyr Leu Phe Ser
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Ala Val Gly Arg Gly Arg Pro Ala Ser Tyr Leu Trp Asp Leu Gly Asp
1490 1495 1500

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Glu Ala Trp Leu Asn Val Thr Val Lys Arg Arg Val Arg Gly Leu Val
1540 1545 1550

gtc aat gca agc cgc acg gtg gtg ccc ctg aat ggg agc gtg agc ttc 4704
Val Asn Ala Ser Arg Thr Val Val Pro Leu Asn Gly Ser Val Ser Phe
1555 1560 1565

agc acg tcc ctg gag gcc ggc agt gat gtg cgc tat tcc tgg gtg ctc 4752
Ser Thr Ser Leu Glu Ala Gly Ser Asp Val Arg Tyr Ser Trp Val Leu
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Cys Asp Arg Cys Thr Pro Ile Pro Gly Gly Pro Thr Ile Ser Tyr Thr
1585 1590 1595 1600

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Phe Arg Ser Val Gly Thr Phe Asn Ile Ile Val Thr Ala Glu Asn Glu
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Val Gly Ser Ala Gln Asp Ser Ile Phe Val Tyr Val Leu Gln Leu Ile
1620 1625 1630

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1635 1640 1645

acg gta cag ctg cag gcc gtg gtt agg gat ggc acc aac gtc tcc tac 4992
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Lys Gly Phe Ser Leu Thr Val Leu Glu Ala Gly Thr Tyr His Val Gln
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Leu Arg Ala Thr Asn Met Leu Glu Ser Ala Trp Ala Asp Cys Thr Met
1700 1705 1710

gac ttc gtg gag cct gtg ggg tgg ctg atg gtg gcc gcc tcc ccg aac 5184
Asp Phe Val Glu Pro Val Gly Trp Leu Met Val Ala Aia Ser Pro Asn
1715 1720 1725

cca gct gcc gtc aac aca agc gtc acc ctc agt gcc gag ctg gct ggt 5232
Pro Ala Ala Val Asn Thr Ser Val Thr Leu Ser Ala Glu Leu Ala Gly
1730 1735 1740

ggc agt ggt gtc gta tac act tgg tcc ttg gag gag ggg ctg agc tgg Gly Ser Gly Val Val Tyr Thr Trp Ser Leu Glu Glu Ser Ser Trp 1745 1750 1755 1760	5280
gag acc tcc gag cca ttt acc acc cat agr ttc ccc aca ccc ggc ctg Glu Thr Ser Glu Pro Phe Thr Thr His Ser Phe Pro Thr Pro Gly Leu 1765 1770 1775	5328
cac ttg gtc acc atg acg gca ggg aac ccc ctg ggc tca gcc aac gcc His Leu Val Thr Met Thr Ala Gly Asn Pro Leu Gly Ser Ala Asn Ala 1780 1785 1790	5376
acc gtg gaa gtg gat gtg cag gtg cct gtg agt ggc ctc aca atc agg Thr Val Glu Val Asp Val Gln Val Pro Val Ser Gly Leu Ser Ile Arg 1795 1800 1805	5424
gcc agc gag ccc gga ggc aca ccc gtg gca gcc ggg tcc tct gtg ccc Ala Ser Glu Pro Gly Ser Phe Val Ala Ala Gly Ser Ser Val Pro 1810 1815 1820	5472
ttt tgg ggg cag ctg gcc acc aat gtg agc tgg tgg tgg gct Phe Trp Gly Gln Leu Ala Thr Gly Thr Asn Val Ser Trp Cys Trp Ala 1825 1830 1835 1840	5520
gtg ccc ggc ggc aca aca aag cgt ggc cct cat gtc acc atc gtc tcc Val Pro Gly Gly Ser Ser Lys Arg Gly Pro His Val Thr Met Val Phe 1845 1850 1855	5568
ccg gat gct ggc acc ttc tcc atc cgg ctc aat gcc tcc aac gca gtc Pro Asp Ala Gly Thr Phe Ser Ile Arg Leu Asn Ala Ser Asn Ala Val 1860 1865 1870	5616
agc tgg gtc tca gcc acc tac aac ctc acg gcg gag gag ccc atc gtg Ser Trp Val Ser Ala Thr Tyr Asn Leu Thr Ala Glu Glu Pro Ile Val 1875 1880 1885	5664
ggc ctg gtg ctg tgg gcc agc aac gtc aag gtg gtc gca ggg ccc gag ctg Gly Leu Val Leu Trp Ala Ser Ser Lys Val Val Ala Pro Gly Gln Leu 1890 1895 1900	5712
gtc cat ttt cag atc ctg ctg gct gcc ggc tca gct gtc acc ttc cgc Val His Phe Gln Ile Leu Ala Ala Gly Ser Ala Val Thr Phe Arg 1905 1910 1915 1920	5760
cta cag gtc ggc ggg gcc aac ccc gag gtg ctc ccc ggg ccc cgt ttc Leu Gln Val Gly Gly Ala Asn Pro Glu Val Leu Pro Gly Pro Arg Phe 1925 1930 1935	5808
tcc cac agc ttc ccc cgc gtc gga gac cac gtg gtg agc gtg cgg ggc Ser His Ser Phe Pro Arg Val Gly Asp His Val Val Ser Val Arg Gly 1940 1945 1950	5856
aaa aac cac gtg agc tgg gcc cag gcg cag gtg cgc atc gtg gtg ctg Lys Asn His Val Ser Trp Ala Gln Ala Gln Val Arg Ile Val Val Leu 1955 1960 1965	5904
gag gcc gtg agt ggg ctg cag gtg ccc aac tgc tgc gag cct ggc atc Glu Ala Val Ser Gly Leu Gln Val Pro Asn Cys Cys Glu Pro Gly Ile 1970 1975 1980	5952
gcc acg ggc act gag agg aac ttc aca gcc cgc gtg cag cgc ggc tct Ala Thr Gly Thr Glu Arg Asn Phe Thr Ala Arg Val Gln Arg Gly Ser 1985 1990 1995 2000	6000
cgg gtc gcc tac gcc tgg tac ttc tcg ctg cag aag gtc cag ggc gac Arg Val Ala Tyr Ala Trp Tyr Phe Ser Leu Gln Lys Val Gln Gly Asp 2005 2010 2015	6048

tcg ctg gtc atc ctg tcg ggg cgc gag gtc acc tac acg ccc gtg gcc	6096
Ser Leu Val Ile Leu Ser Gly Arg Asp Val Thr Tyr Thr Pro Val Ala	
2020 2025 2030	
gca ggg ctg ttg gag atc cag gtg cgc gac ttc aac gct ctg ggc agt	6144
Ala Gly Leu Leu Glu Ile Gln Val Arg Ala Phe Asn Ala Leu Gly Ser	
2035 2040 2045	
gag aac cgc acg ctg gtc ctg gag gtt cag gag gca gtc cag tat gtg	6192
Glu Asn Arg Thr Leu Val Leu Glu Val Gln Asp Ala Val Gln Tyr Val	
2050 2055 2060	
gcc ctg cag acg ggc ccc tgc ttc acc aac cgc tcc ggc cag ttt gag	6240
Ala Leu Gln Ser Gly Pro Cys Phe Thr Asn Arg Ser Ala Gln Phe Glu	
2065 2070 2075 2080	
gcc gcc acc acg ccc acg ccc cgg cgt gtg gcc tac cac tgg gac ttt	6283
Ala Ala Thr Ser Pro Ser Pro Arg Arg Val Ala Tyr His Trp Asp Phe	
2085 2090 2095	
ggg gat ggg tcc cca ggg cag gag aca gat gag ccc agg gcc gag cac	6336
Gly Asp Gly Ser Pro Gly Gln Asp Thr Asp Glu Pro Arg Ala Glu His	
2100 2105 2110	
tcc tac ctg agg cct ggg gac tac cgc gtg cag gtg aac gcc tcc aac	6384
Ser Tyr Leu Arg Pro Gly Asp Tyr Arg Val Gln Val Asn Ala Ser Asn	
2115 2120 2125	
ctg gtg agc ttc ttc gtg gcc cag gtc acc gtc cag gtg ctg	6432
Leu Val Ser Phe Phe Val Ala Gln Ala Thr Val Thr Val Gln Val Leu	
2130 2135 2140	
gcc tgc cgg gag ccc gag gtg gac gtg gtc ctg ccc ctg cag gtg ctg	6480
Ala Cys Arg Glu Pro Glu Val Asp Val Val Leu Pro Leu Gln Val Leu	
2145 2150 2155 2160	
atg cgg cga tca cag cgc aac tac ttg gag gcc cac gtt gac ctg cgc	6528
Met Arg Arg Ser Gln Arg Asn Tyr Leu Glu Ala His Val Asp Leu Arg	
2165 2170 2175	
gac tgc gtc acc tac cag act gag tac cgc tgg gag gtg tat cgc acc	6576
Asp Cys Val Thr Tyr Gln Thr Glu Tyr Arg Trp Glu Val Tyr Arg Thr	
2180 2185 2190	
gcc agc tgc cag cgg ccc ggg cgc cca gcg cgt gtg gcc ctg ccc ggc	6624
Ala Ser Cys Gln Arg Pro Gly Arg Pro Ala Arg Val Ala Leu Pro Gly	
2195 2200 2205	
gtg gag gtg agc cgg cct cgg ctg gtg ctg ccc cgg ctg gcg ctg cct	6672
Val Asp Val Ser Arg Pro Arg Leu Val Leu Pro Arg Leu Ala Leu Pro	
2210 2215 2220	
gtg ggg cac tac tgc ttt gtg ttt gtc gtg tca ttt ggg gag acg cca	6720
Val Gly His Tyr Cys Phe Val Phe Val Ser Phe Gly Asp Thr Pro	
2225 2230 2235 2240	
ctg aca cag agc atc cag gcc aat gtg acg gtg gcc ccc gag cgc ctg	6768
Leu Thr Gln Ser Ile Gln Ala Asn Val Thr Val Ala Pro Glu Arg Leu	
2245 2250 2255	
gtg ccc atc att gag ggt ggc tca tac cgc gtg tgg tca gag aca cgg	6816
Val Pro Ile Ile Glu Gly Ser Tyr Arg Val Trp Ser Asp Thr Arg	
2260 2265 2270	
gac ctg gtg ctg gat ggg agc gag tcc tac gag ccc aac ctg gag gac	6864
Asp Leu Val Leu Asp Gly Ser Glu Ser Tyr Asp Pro Asn Leu Glu Asp	

2275	2280	2285	
ggc gac cag acg ccg ctc agt ttc cac tgg gcc tgt gtc gct tgg aca Gly Asp Gln Thr Pro Leu Ser Phe His Trp Ala Cys Val Ala Ser Thr 2290	2295	2300	6912
caj agg gag gct ggc ggg tgt gcg ctg aac ttt ggg ccc cgc ggg agc Gln Arg Glu Ala Gly Gly Cys Ala Leu Asn Phe Gly Pro Arg Gly Ser 2305	2310	2315	6960
agg acg gtc acc att cca cgg gag cgg ctg ggg gct ggc gtc gag tac Ser Thr Val Thr Ile Pro Arg Glu Arg Leu Ala Ala Gly Val Glu Tyr 2325	2330	2335	7008
acc ttc aca ctg acc gtc tgg aag gcc ggc cgg aac gag gag gac acc Thr Phe Ser Leu Thr Val Trp Lys Ala Gly Arg Lys Glu Glu Ala Thr 2340	2345	2350	7056
aac caa acg gtg ctg atc cgg aat ggc cgg gtc ccc att gtc tcc tgg Asn Gln Thr Val Leu Ile Arg Ser Gly Arg Val Pro Ile Val Ser Leu 2355	2360	2365	7104
gag tgt gtg tcc tgg aag gca cag gcc gtc tac gaa gtg aac cgc aac Glu Cys Val Ser Cys Lys Ala Gln Ala Val Tyr Glu Val Ser Arg Ser 2370	2375	2380	7152
tcc tac gtg tac tgg gag ggc cgc tgg ctc aat tyc aac aac gtc tcc Ser Tyr Val Tyr Leu Glu Gly Arg Cys Leu Asn Cys Ser Ser Gly Ser 2385	2390	2395	7200
aag cga ggg cgg tgg gct gca cgt aac ttc aac aac aac ctc gtg Lys Arg Gly Arg Trp Ala Ala Arg Thr Phe Ser Asn Lys Thr Leu Val 2405	2410	2415	7248
ctg gat gag acc acc aca tcc acg ggc aat gca ggc atc cga ctc gtg Leu Asp Glu Thr Thr Ser Thr Gly Ser Ala Gly Met Arg Leu Val 2420	2425	2430	7296
ctg cgg cgg ggc gtg ctg cgg gac ggc gag gga tac acc ttc aac ctc Leu Arg Arg Gly Val Leu Arg Asp Gly Glu Gly Tyr Thr Phe Thr Leu 2435	2440	2445	7344
acg gtg ctg ggc cgc tct ggc gag gag ggc tgc gcc tcc atc cgc Thr Val Leu Gly Arg Ser Gly Glu Glu Gly Cys Ala Ser Ile Arg 2450	2455	2460	7392
ctg tcc ccc aac cgc ccc ctg ggg ggc tct tgc cgc ctc ttc cca Leu Ser Pro Asn Arg Pro Leu Gly Gly Ser Cys Arg Leu Phe Pro 2465	2470	2475	7440
ctg ggc gct gtg cac gcc ctc acc acc aac gtg cac ttc gaa tgc acg Leu Gly Ala Val His Ala Leu Thr Thr Lys Val His Phe Glu Cys Thr 2485	2490	2495	7488
ggc tgg cat gac gcg gag gat gct ggc gcc ccc ctg gtg tac gcc ctc Gly Trp His Asp Ala Glu Asp Ala Gly Ala Pro Leu Val Tyr Ala Leu 2500	2505	2510	7536
ctg ctg cgg cgc tgt cgc cag ggc cac tgc gag gag ttc tgt gtc tac Leu Leu Arg Arg Cys Arg Gln Gly His Cys Glu Glu Phe Cys Val Tyr 2515	2520	2525	7584
aag ggc aac ctc tcc agc tac gga gcc gtc ccc ccc ggt ttc agg Lys Gly Ser Leu Ser Ser Tyr Gly Ala Val Leu Pro Pro Gly Phe Arg 2530	2535	2540	7632
cca cac ttc gag gtg ggc ctg gcc gtc gtc gtc cag gac cag ctg gga			7680

Pro His Phe Glu Val Gly Leu Ala Val Val Gln Asp Gln Leu Gly				
2545	2550	2555	2560	
gcc gct gtg gtc gcc ctc aac agg tct ttg gcc atc acc ctc cca gag				7728
Ala Ala Val Val Ala Leu Asn Arg Ser Leu Ala Ile Thr Leu Pro Glu				
2565	2570	2575		
ccc aac ggc aac gca acg ggg ctc aca gtc tgg ctg cac ggg ctc acc				7776
Pro Asn Gly Ser Ala Thr Gly Leu Thr Val Trp Leu His Gly Leu Thr				
2580	2585	2590		
gtc aat gtg ctc cca ggg ctc ctt cgg cag gcc gat ccc cag cac gtc				7824
Ala Ser Val Leu Pro Gly Leu Leu Arg Gln Ala Asp Pro Gln His Val				
2595	2600	2605		
atc gag tac tcc ttg gcc ctc gtc acc gtg ctc aac gag tac gag cgg				7872
Ile Glu Tyr Ser Leu Ala Leu Val Thr Val Leu Asn Glu Tyr Glu Arg				
2610	2615	2620		
gcc ctc gag gtc gcg gca gag ccc aag cac gag cgg cag ccc gca gca				7920
Ala Leu Asp Val Ala Ala Pro Lys His Glu Arg Gln His Arg Ala				
2625	2630	2635	2640	
cag ata cgc aag aac atc acg gag act ctc gtg tcc ctc aag gtc cac				7968
Gln Ile Arg Lys Asn Ile Thr Glu Thr Leu Val Ser Leu Arg Val His				
2645	2650	2655		
act gtg gat gac atc cag cag atc gct gct gcg ctc gcc cag tcc atg				8016
Thr Val Asp Asp Ile Gln Gln Ile Ala Ala Leu Ala Gln Cys Met				
2660	2665	2670		
ggg ccc aac agg gag ctc gta tgc cgc tgc ctg aag cag ccc ctc				8064
Gly Pro Ser Arg Glu Leu Val Cys Arg Ser Cys Leu Lys Gln Thr Leu				
2675	2680	2685		
cac aag ctc gag gcc atg atg ctc atc ctc cag gca gag acc acc gcg				8112
His Lys Leu Glu Ala Met Met Leu Ile Leu Gln Ala Glu Thr Thr Ala				
2690	2695	2700		
ggc acc gtg acg ccc acc gcc atc gga gac agc atc ctc aac atc aca				8160
Gly Thr Val Thr Pro Thr Ala Ile Gly Asp Ser Ile Leu Asn Ile Thr				
2705	2710	2715	2720	
gga gac ctc atc cac ctc gcc agc tcc gac gtg cgg gca cca cag ccc				8208
Gly Asp Leu Ile His Leu Ala Ser Ser Asp Val Arg Ala Pro Gln Pro				
2725	2730	2735		
tca gag ctc gga gcc gag tca cca tct cgg atg gtg gcg tcc cag gcc				8256
Ser Glu Leu Gly Ala Glu Ser Pro Ser Arg Met Val Ala Ser Gln Ala				
2740	2745	2750		
tac aac ctc acc tct gcc ctc atg cgc atc ctc atg cgc tcc cgc gtg				8304
Tyr Asn Leu Thr Ser Ala Leu Met Arg Ile Leu Met Arg Ser Arg Val				
2755	2760	2765		
ctc aac gag gag ccc ctc acg ctc gcg ggc gag qag atc gtg gcc cag				8352
Leu Asn Glu Glu Pro Leu Thr Leu Ala Gly Glu Glu Ile Val Ala Gln				
2770	2775	2780		
ggc aag cgc tcc gac ccc cgg agc ctc tcc tat ggc ggc gcc cca				8400
Gly Lys Arg Ser Asp Pro Arg Ser Leu Leu Cys Tyr Gly Ala Pro				
2785	2790	2795	2800	
ggc cct ggc tcc cac ttc tcc atc ccc gag gct ttc agc ggg gcc ctc				8448
Gly Pro Gly Cys His Phe Ser Ile Pro Glu Ala Phe Ser Gly Ala Leu				
2805	2810	2815		

gcc aac ctc agt gag gtg gtg cag ctc atc ttt ctg gtg gac tcc aat		8496	
Ala Asn Leu Ser Asp Val Val Gln Leu Ile Phe Leu Val Asp Ser Asn			
2820	2825	2830	
ccc ttt ccc ttt ggg tat atc aac tac acc gtc tcc acc aag gtg		8544	
Pro Phe Pro Phe Gly Tyr Ile Ser Asn Tyr Thr Val Ser Thr Lys Val			
2835	2840	2845	
gcc tcc tct gca ttc cag aca cag gcc ggc gcc cag atc ccc acc gag		8592	
Ala Ser Met Ala Phe Gln Thr Gln Ala Gly Ala Gln Ile Pro Ile Glu			
2850	2855	2860	
cgg ctg gcc tca gag cgc gcc acc acc gtg aag gtg ccc aac aac tcc		8640	
Arg Leu Ala Ser Glu Arg Ala Ile Thr Val Lys Val Pro Asn Asn Ser			
2865	2870	2875	2880
gac tgg gct gcc cgg ggc dac cgc agc tcc gcc aac tcc gcc aac tcc		8684	
Asp Trp Ala Ala Arg Gly His Arg Ser Ser Ala Asn Ser Ala Asn Ser			
2885	2890	2895	
gtt gtg gtc cag ccc cag gcc tcc gtc ggt gct gtg gtc acc ctg gac		8736	
Val Val Val Gln Pro Gln Ala Ser Val Gly Ala Val Val Thr Leu Asp			
2900	2905	2910	
agg aac aac cct gcg gcc ggg ctg cat ctg cag ctc aac tat acc ctg		8784	
Ser Ser Asn Pro Ala Ala Gly Leu His Leu Gln Leu Asn Tyr Thr Leu			
2915	2920	2925	
ctg gac ggc cac tac ctg tct gag gaa cct gag ccc tac ctg gca gtc		8832	
Leu Asp Gly His Tyr Leu Ser Glu Glu Pro Glu Pro Tyr Leu Ala Val			
2930	2935	2940	
tac cta cac tcc gag ccc cgg ccc aat gag cac aac tcc tcc gct aac		8880	
Tyr Leu His Ser Glu Pro Arg Pro Asn Glu His Asn Cys Ser Ala Ser			
2945	2950	2955	2960
agg agg atc cgc cca gag tca ctc cag ggt gct gac cac cgg ccc tac		8928	
Arg Arg Ile Arg Pro Glu Ser Leu Gln Gly Ala Asp His Arg Pro Tyr			
2965	2970	2975	
acc ttc ttc att tcc ccc ggg agc aga gac cca gcg ggg agt tac cat		8976	
Thr Phe Ile Ser Pro Gly Ser Arg Asp Pro Ala Gly Ser Tyr His			
2980	2985	2990	
ctg aac ctc tcc agc cac ttc cgc tgg tcg gcg ctg cag gtg tcc gtg		9024	
Leu Asn Leu Ser Ser His Phe Arg Trp Ser Ala Leu Gln Val Ser Val			
2995	3000	3005	
ggc ctg tac acg tcc ctg tgc cag tac ttc agc gag gag gac atg gtg		9072	
Gly Leu Tyr Thr Ser Leu Cys Gln Tyr Phe Ser Glu Glu Asp Met Val			
3010	3015	3020	
tgg cgg aca gag ggg ctg ctg ccc ctg gag gag acc tcc ccc cgc cag		9120	
Trp Arg Thr Glu Gly Leu Leu Pro Leu Glu Thr Ser Pro Arg Gln			
3025	3030	3035	3040
gcc gtc tcc acc cgc cac ctc acc gcc ttc ggc gcc agc ctc ttc		9168	
Ala Val Cys Leu Thr Arg His Leu Thr Ala Phe Gly Ala Ser Leu Phe			
3045	3050	3055	
gtg ccc cca agc cat gtc cgc ttt gtg ttt cct gag ccc aca gcg gat		9216	
Val Pro Pro Ser His Val Arg Phe Val Phe Pro Glu Pro Thr Ala Asp			
3060	3065	3070	
gta aac tac atc gtc atg ctg aca tgt gct gtg tgc ctg gtg acc tac		9264	
Val Asn Tyr Ile Val Met Leu Thr Cys Ala Val Cys Leu Val Thr Tyr			
3075	3080	3085	

3345	3350	3355	3360	
ggg cag cag gtg ctg gac atc gac agc tgc ctg gac tcg tcc gtg ctg Gly Gln Gln Val Leu Asp Ile Asp Ser Cys Leu Asp Ser Ser Val Leu 3365		3370		10128
		3375		
gac agc tcc ttc ctc acg ttc tca ggc ctc cac gct gag cag gcc ttt Asp Ser Ser Phe Leu Thr Phe Ser Gly Leu His Ala Glu Gln Ala Phe 3380	3385		3390	10176
gtt gga cag atg aag agt gac ttg ttt ctg gat gat tct aag agt ctg Val Gly Gln Met Lys Ser Asp Leu Phe Leu Asp Asp Ser Lys Ser Leu 3395	3400	3405		10224
gtg tgc tgg ccc tcc ggc gag gga acg ctc agt tgg ccg gac ctg ctc Val Cys Trp Pro Ser Gly Glu Gly Thr Leu Ser Trp Pro Asp Leu Leu 3410	3415	3420		10272
agt gac ccc tcc att gtg ggt agc aat ctg cgg cag ctg gca cgg ggc Ser Asp Pro Ser Ile Val Gly Ser Asn Leu Arg Gln Leu Ala Arg Gly 3425	3430	3435	3440	10320
cag gcg ggc cat ggg ctg ggc cca gag gag gac ggc ttc tcc ctg gcc Gln Ala Gly His Gly Leu Gly Pro Glu Glu Asp Gly Phe Ser Leu Ala 3445	3450	3455		10368
agc ccc tac tcg cct gcc aaa tcc ttc tca gca tca gat gaa gac ctg Ser Pro Tyr Ser Pro Ala Lys Ser Phe Ser Ala Ser Asp Glu Asp Leu 3460	3465	3470		10416
atc cag cag gtc ctt gcc gag ggg gtc agc agc cca gcc cct acc caa Ile Gln Gln Val Leu Ala Glu Gly Val Ser Ser Pro Ala Pro Thr Gln 3475	3480	3485		10464
gac acc cac atg gaa acg gac ctg ctc agc agc ctg tcc agc act cct Asp Thr His Met Glu Thr Asp Leu Leu Ser Ser Leu Ser Ser Thr Pro 3490	3495	3500		10512
ggg gag aag aca gag acg ctg gcg ctg cag agg ctg ggg gag ctg ggg Gly Glu Lys Thr Glu Thr Ala Leu Gln Arg Leu Gly Glu Leu Gly 3505	3510	3515	3520	10560
cca ccc agc cca ggc ctg aac tgg jaa cag ccc cag gca gcg agg ctg Pro Pro Ser Pro Gly Leu Asn Trp Glu Gln Pro Gln Ala Ala Arg Leu 3525	3530	3535		10608
tcc agg aca gga ctg gtg gag ggt ctg cgg aag cgc ctg ctg ccg gcc Ser Arg Thr Gly Leu Val Glu Gly Leu Arg Lys Arg Leu Leu Pro Ala 3540	3545	3550		10656
tgg tgt gcc tcc ctg gcc cac ggg ctc agc ctg ctc ctg gtg gct gtg Trp Cys Ala Ser Leu Ala His Gly Leu Ser Leu Leu Val Ala Val 3555	3560	3565		10704
gct gtg gct gtc tca ggg tgg gtg ggt gcg agc ttc ccc ccg ggc gtg Ala Val Ala Val Ser Gly Trp Val Gly Ala Ser Phe Pro Pro Gly Val 3570	3575	3580		10752
agt gtt gcg tgg ctc ctg tcc agc agc gcc agc ttc ctg gcc tca ttc Ser Val Ala Trp Leu Leu Ser Ser Ala Ser Phe Leu Ala Ser Phe 3585	3590	3595	3600	10800
ctc ggc tgg gag cca ctg aag gtc ttg ctg gaa gcc ctg tac ttc tca Leu Gly Trp Glu Pro Leu Lys Val Leu Leu Glu Ala Leu Tyr Phe Ser 3605	3610	3615		10848
ctg gtg gcc aag cgg ctg cac ccg gat gaa gat gac acc ctg gta gag				10896

Leu Val Ala Lys Arg Leu His Pro Asp Glu Asp Asp Thr Leu Val Glu			
3620	3625	3630	
agc ccg gct gtg acc cct gtg agc gca aat gtc gtg ccc cgc gta cgg cca			10944
Ser Pro Ala Val Thr Pro Val Ser Ala Arg Val Pro Arg Val Arg Pro			
3635	3640	3645	
ccc cac ggc ttt gca ctc ttc ctg gcc aag gaa gaa gcc cgc aag gtc			10992
Pro His Gly Phe Ala Leu Phe Leu Ala Lys Glu Glu Ala Arg Lys Val			
3650	3655	3660	
aag agg cta cat ggc atg ctg cgg agc ctc ctg gtg tac atg ctt ttt			11040
Lys Arg Leu His Gly Met Leu Arg Ser Leu Leu Val Tyr Met Leu Phe			
3665	3670	3675	3680
ctg ctg gtg acc ctg ctg gcc agc tat ggg gat gcc tca tgc cat ggg			11088
Leu Leu Val Thr Leu Ala Ser Tyr Gly Asp Ala Ser Cys His Gly			
3685	3690	3695	
cac gcc tac aat ctg ctt aag gcc atc aag gag gag ctg sac agc cgg			11136
His Ala Tyr Arg Leu Gln Ser Ala Ile Lys Gln Glu Leu His Ser Arg			
3700	3705	3710	
gcc ttc ctg gcc atc acg cgg tct gag gag ctc tgg cca tgg atg gcc			11184
Ala Phe Leu Ala Ile Thr Arg Ser Glu Glu Leu Trp Pro Trp Met Ala			
3715	3720	3725	
cac gtg ctg ctg ccc tac gtc cac ggg aac cag tcc aat cca gag ctg			11232
His Val Leu Leu Pro Tyr Val His Gly Asn Gln Ser Ser Pro Glu Leu			
3730	3735	3740	
ggg ccc cca cgg ctg cgg cag gtg cgg ctg cag gaa gca ctc tac cca			11280
Gly Pro Pro Arg Leu Arg Gln Val Arg Leu Gln Glu Ala Leu Tyr Pro			
3745	3750	3755	3760
gac cct ccc gcc ccc agg gtc cac acg tgc tgg gcc gca gga ggc ttc			11328
Asp Pro Pro Gly Pro Arg Val His Thr Cys Ser Ala Ala Gly Gly Phe			
3765	3770	3775	
agc acc aac gat tac gac gtt ggc tgg gag aat cct cac aat ggc tcc			11376
Ser Thr Ser Asp Tyr Asp Val Gly Trp Glu Ser Pro His Asn Gly Ser			
3780	3785	3790	
ggg acg tgg gcc tat tca gcg ccg gat ctg ctg ggg gca tgg tcc tgg			11424
Gly Thr Trp Ala Tyr Ser Ala Pro Asp Leu Leu Gly Ala Trp Ser Trp			
3795	3800	3805	
ggc tcc tgt gcc gtg tat gac aac ggg ggc tac gtg cag gag ctg ggc			11472
Gly Ser Cys Ala Val Tyr Asp Ser Gly Gly Tyr Val Gln Glu Leu Gly			
3810	3815	3820	
ctg agc ctg gag gag agc cgc gac cgg ctg cgc ttc ctg cag ctg cac			11520
Leu Ser Leu Glu Ser Arg Asp Arg Leu Arg Phe Leu Gln Leu His			
3825	3830	3835	3840
aac tgg ctg gac aac agg agc cgc gct gtg ttc ctg gag ctc acg cgc			11568
Asn Trp Leu Asp Asn Arg Ser Arg Ala Val Phe Leu Glu Leu Thr Arg			
3845	3850	3855	
tac agc ccg gcc gtg ggg ctg cac gcc gcc gtc acg ctg cgc ctc gag			11616
Tyr Ser Pro Ala Val Gly Leu His Ala Ala Val Thr Leu Arg Leu Glu			
3860	3865	3870	
ttc ccg cgc gcc ggc cgc gcc ctg gcc gcc ctc acg gtc cgc ccc ttt			11664
Phe Pro Ala Ala Gly Arg Ala Leu Ala Ala Leu Ser Val Arg Pro Phe			
3875	3880	3885	

ggc tgc cgc ctc agc ggc ctc tgg ctg ctc ctc acc tgg Ala Leu Arg Arg Leu Ser Ala Gly Leu Ser Leu Pro Leu Leu Thr Ser	3890	3895	3900	11712	
gtg tgc ctg ctg ttc gcc gtg cac ttc gcc gtg gcc gag gcc cgt Val Cys Leu Leu Leu Phe Ala Val His Phe Ala Val Ala Glu Ala Arg	3905	3910	3915	3920	11760
act tgg cac agg jaa ggg cgc tgg cgc gtg ctg tgg ctc gga gcc tgg Thr Trp His Arg Glu Gly Arg Trp Arg Val Leu Arg Leu Gly Ala Trp	3925	3930	3935		11803
ggc cgg tgg ctg ctg gtg ggc ctg acc ggc gcc acc gca ctg gta cgc Ala Arg Trp Leu Leu Val Ala Leu Thr Ala Ala Thr Ala Leu Val Arg	3940	3945	3950		11856
ctc gcc cag ctg ggt gcc gct gag cgc cag tgg acc cgt ttc gtg cgc Leu Ala Glu Leu Gly Ala Ala Asp Arg Gln Trp Thr Arg Phe Val Arg	3955	3960	3965		11904
ggc cgc cgc cgc ttc act acc ttc gag cag gtg ggc cag gtg agg Gly Arg Pro Arg Arg Phe Thr Ser Phe Asp Gln Val Ala His Val Ser	3970	3975	3980		11952
tcc gca gcc cgt ggc ctg ggc tcc ctg ctc ttc ctg ctt ttg gtc Ser Ala Ala Arg Gly Leu Ala Ala Ser Leu Leu Phe Leu Leu Val	3985	3990	3995	4000	12000
aag gct ggc cag cac gta cgc ttc gtg cgc cag tgg tcc gtc ttt ggc Lys Ala Ala Gln His Val Arg Phe Val Arg Gln Trp Ser Val Phe Gly	4005	4010	4015		12048
aag aca tta tgc cga gct ctg cca gag ctc ctg ggg gtc acc ttg ggc Dys Thr Leu Cys Arg Ala Leu Pro Glu Leu Leu Gly Val Thr Leu Gly	4020	4025	4030		12096
ctg gtg gtc ggg gta gcc tac gcc cag ctg ggc acc ctg ctc gtg Leu Val Val Gly Val Ala Tyr Ala Gln Leu Ala Ile Leu Leu Val	4035	4040	4045		12144
tct tcc tgt gtg gag tcc ctc tgg agc gtg gcc cag gcc ctg ttg gtg Ser Ser Cys Val Asp Ser Leu Trp Ser Val Ala Gln Ala Leu Leu Val	4050	4055	4060		12192
ctg tgc cct ggg act ggg ctc tct acc ctg tgt cct gcc gag tcc tgg Leu Cys Pro Gly Thr Gly Leu Ser Thr Leu Cys Pro Ala Glu Ser Trp	4065	4070	4075	4080	12240
cac ctg tca ccc ctg ctg tgt gtg ggg ctc tgg gca ctg cgg ctg tgg His Leu Ser Pro Leu Leu Cys Val Gly Leu Trp Ala Leu Arg Leu Trp	4085	4090	4095		12288
ggc gcc cta cgg ctg ggg gct gtt att ctc cgc tgg cgc tac cac gcc Gly Ala Leu Arg Leu Gly Ala Val Ile Leu Arg Trp Arg Tyr His Ala	4100	4105	4110		12336
ttg cgt gga gag ctg tac cgg ccg gcc tgg gag ccc cag gag tac gag Leu Arg Gly Glu Leu Tyr Arg Pro Ala Trp Glu Pro Gln Asp Tyr Glu	4115	4120	4125		12384
atg gtg gag ttg ttc ctg cgc agg ctg cgc ctc tgg atg ggc ctc agc Met Val Glu Leu Phe Leu Arg Arg Leu Arg Leu Trp Met Gly Leu Ser	4130	4135	4140		12432
aag gtc aag gag ttc cgc cac aaa gtc cgc ttt gaa ggg atg gag ccg Lys Val Lys Glu Phe Arg His Lys Val Arg Phe Glu Gly Met Glu Pro	4145	4150	4155	4160	12480

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Leu Pro Ser Arg Ser Ser Arg Gly Ser Lys Val Ser Pro Asp Val Pro
4165 4170 4175

cca ccc aac gct ggc tcc gat gcc tcc cac ccc tcc aac tcc tcc aac 12576
Pro Pro Ser Ala Gly Ser Asp Ala Ser His Pro Ser Thr Ser Ser
4180 4185 4190

cag ctg gat ggg ctg aac gtg aac ctg ggc cgg ctg ggg aca aac tgc 12624
Gln Leu Asp Gly Leu Ser Val Ser Leu Gly Arg Leu Gly Thr Arg Cys
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gag cct gag ccc tcc cgg ctc caa ggc gtg ttc gag gcc ctg ctc acc 12672
Glu Pro Glu Pro Ser Arg Leu Gln Ala Val Phe Glu Ala Leu Leu Thr
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Gly Ser Ser Arg Gly Pro Ser Pro Gly Leu Arg Pro Ala Leu Pro Ser
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Cys Glu Pro Pro Cys Leu Cys Gly Pro Ala Pro Gly Ala Ala Cys Arg
35 40 45

Val Asn Cys Ser Gly Arg Gly Leu Arg Thr Leu Gly Pro Ala Leu Arg
50 55 60

Ile Pro Ala Asp Ala Thr Glu Leu Asp Val Ser His Asn Leu Leu Arg
65 70 75 80

Ala Leu Asp Val Gly Leu Leu Ala Asn Leu Ser Ala Leu Ala Glu Leu
85 90 95

Asp Ile Ser Asn Asn Lys Ile Ser Thr Leu Glu Glu Gly Ile Phe Ala
100 105 110

Asn Leu Phe Asn Leu Ser Glu Ile Asn Leu Ser Gly Asn Pro Phe Glu
115 120 125

Cys Asp Cys Gly Leu Ala Trp Leu Pro Gln Trp Ala Glu Glu Gln Gln
 130 135 140
 Val Arg Val Val Gln Pro Glu Ala Ala Thr Cys Ala Gly Pro Gly Ser
 145 150 155 160
 Leu Ala Gly Gln Pro Leu Leu Gly Ile Pro Leu Leu Asp Ser Gly Cys
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 Gly Glu Glu Tyr Val Ala Cys Leu Pro Asp Asn Ser Ser Gly Thr Val
 180 185 190
 Ala Ala Val Ser Phe Ser Ala Ala His Glu Gly Leu Leu Gln Pro Glu
 195 200 205
 Ala Cys Ser Ala Phe Cys Phe Ser Thr Gly Gln Gly Leu Ala Ala Leu
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 Ser Glu Gln Gly Trp Cys Leu Cys Gly Ala Ala Gln Pro Ser Ser Ala
 225 230 235 240
 Ser Phe Ala Cys Leu Ser Leu Cys Ser Gly Pro Pro Ala Pro Pro Ala
 245 250 255
 Pro Thr Cys Arg Gly Pro Thr Leu Leu Gln His Val Phe Pro Ala Ser
 260 265 270
 Pro Gly Ala Thr Leu Val Gly Pro His Gly Pro Leu Ala Ser Gly Gln
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 Leu Ala Ala Phe His Ile Ala Ala Pro Leu Pro Val Thr Asp Thr Arg
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 Trp Asp Phe Gly Asp Gly Ser Ala Glu Val Asp Ala Ala Gly Pro Ala
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 Ala Ser His Arg Tyr Val Leu Pro Gly Arg Tyr His Val Thr Ala Val
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 Glu Ala Ala Pro Ala Ala Leu Glu Leu Val Cys Pro Ser Ser Val Gln
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 Ser Asp Glu Ser Leu Asp Leu Ser Ile Gln Asn Arg Gly Gly Ser Gly
 370 375 380
 Leu Glu Ala Ala Tyr Ser Ile Val Ala Leu Gly Glu Glu Pro Ala Arg
 385 390 395 400
 Ala Val His Pro Leu Cys Pro Ser Asp Thr Glu Ile Phe Pro Gly Asn
 405 410 415
 Gly His Cys Tyr Arg Leu Val Val Glu Lys Ala Ala Trp Leu Gln Ala
 420 425 430
 Gln Glu Gln Cys Gln Ala Trp Ala Gly Ala Ala Leu Ala Met Val Asp
 435 440 445
 Ser Pro Ala Val Gln Arg Phe Leu Val Ser Arg Val Thr Arg Ser Leu
 450 455 460
 Asp Val Trp Ile Gly Phe Ser Thr Val Gln Gly Val Glu Val Gly Pro
 465 470 475 480

Ala Pro Gln Gly Glu Ala Phe Ser Leu Glu Ser Cys Gln Asn Trp Leu
 435 490 495
 Pro Gly Glu Pro His Pro Ala Thr Ala Gln His Cys Val Arg Leu Gly
 500 505 510
 Pro Thr Gly Trp Cys Asn Thr Asp Leu Cys Ser Ala Pro His Ser Tyr
 515 520 525
 Val Cys Glu Leu Gln Pro Gly Gly Pro Val Gln Asp Ala Glu Asn Leu
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 Leu Val Gly Ala Pro Ser Gly Asp Leu Gln Gly Pro Leu Thr Pro Leu
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 Ala Gln Gln Asp Gly Leu Ser Ala Pro His Glu Pro Val Glu Val Met
 565 570 575
 Val Phe Pro Gly Leu Arg Leu Ser Arg Glu Ala Phe Leu Thr Thr Ala
 580 585 590
 Glu Phe Gly Thr Gln Glu Leu Arg Arg Pro Ala Gln Leu Arg Leu Gln
 595 600 605
 Val Tyr Arg Leu Leu Ser Thr Ala Gly Thr Pro Glu Asn Gly Ser Glu
 610 615 620
 Pro Glu Ser Arg Ser Pro Asp Asn Arg Thr Gln Leu Ala Pro Ala Cys
 625 630 635 640
 Met Pro Gly Gly Arg Trp Cys Pro Gly Ala Asn Ile Cys Leu Pro Leu
 645 650 655
 Asp Ala Ser Cys His Pro Gln Ala Cys Ala Asn Gly Cys Thr Ser Gly
 660 665 670
 Pro Gly Leu Pro Gly Ala Pro Tyr Ala Leu Trp Arg Glu Phe Leu Phe
 675 680 685
 Ser Val Pro Ala Gly Pro Pro Ala Gln Tyr Ser Val Thr Leu His Gly
 690 695 700
 Gln Asp Val Leu Met Leu Pro Gly Asp Leu Val Gly Leu Gln His Asp
 705 710 715 720
 Ala Gly Pro Gly Ala Leu Leu His Cys Ser Pro Ala Pro Gly His Pro
 725 730 735
 Gly Pro Arg Ala Pro Tyr Leu Ser Ala Asn Ala Ser Ser Trp Leu Pro
 740 745 750
 His Leu Pro Ala Gln Leu Glu Gly Thr Trp Gly Cys Pro Ala Cys Ala
 755 760 765
 Leu Arg Leu Leu Ala Gln Arg Glu Gln Leu Thr Val Leu Leu Gly Leu
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 Arg Pro Asn Pro Gly Leu Arg Leu Pro Gly Arg Tyr Glu Val Arg Ala
 785 790 795 800
 Glu Val Gly Asn Gly Val Ser Arg His Asn Leu Ser Cys Ser Phe Asp
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 Val Val Ser Pro Val Ala Gly Leu Arg Val Ile Tyr Pro Ala Pro Arg
 820 825 830
 Asp Gly Arg Leu Tyr Val Pro Thr Asn Gly Ser Ala Leu Val Leu Gln

835 840 845

Val Asp Ser Gly Ala Asn Ala Thr Ala Thr Ala Arg Trp Pro Gly Gly
 850 855 860

Ser Leu Ser Ala Arg Phe Glu Asn Val Cys Pro Ala Leu Val Ala Thr
 865 870 875 880

Phe Val Pro Ala Cys Pro Trp Glu Thr Asn Asp Thr Leu Phe Ser Val
 885 890 895

Val Ala Leu Pro Trp Leu Ser Glu Gly Glu His Val Val Asp Val Val
 900 905 910

Val Glu Asn Ser Ala Ser Arg Ala Asn Leu Ser Leu Arg Val Thr Ala
 915 920 925

Glu Glu Pro Ile Cys Gly Leu Arg Ala Thr Pro Ser Pro Glu Ala Arg
 930 935 940

Val Leu Gln Gly Val Leu Val Arg Tyr Ser Pro Val Val Gln Ala Gly
 945 950 955 960

Ser Asp Met Val Phe Arg Trp Thr Ile Asn Asp Lys Gln Ser Leu Thr
 965 970 975

Phe Gln Asn Val Val Phe Asn Val Ile Tyr Gln Ser Ala Ala Val Phe
 980 985 990

Lys Leu Ser Leu Thr Ala Ser Asn His Val Ser Asn Val Thr Val Asn
 995 1000 1005

Tyr Asn Val Thr Val Glu Arg Met Asn Arg Met Gln Gly Leu Gln Val
 1010 1015 1020

Ser Thr Val Pro Ala Val Leu Ser Pro Asn Ala Thr Leu Ala Leu Thr
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Ala Gly Val Leu Val Asp Ser Ala Val Glu Val Ala Phe Leu Trp Thr
 1045 1050 1055

Phe Gly Asp Gly Glu Gln Ala Leu His Gln Phe Gln Pro Pro Tyr Asn
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Glu Ser Phe Pro Val Pro Asp Pro Ser Val Ala Gln Val Leu Val Glu
 1075 1080 1085

His Asn Val Thr His Thr Tyr Ala Ala Pro Gly Glu Tyr Leu Leu Thr
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Val Leu Ala Ser Asn Ala Phe Glu Asn Leu Thr Gln Gln Val Pro Val
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Ser Val Arg Ala Ser Leu Pro Ser Val Ala Val Gly Val Ser Asp Gly
 1125 1130 1135

Val Leu Val Ala Gly Arg Pro Val Thr Phe Tyr Pro His Pro Leu Pro
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Ser Pro Gly Gly Val Leu Tyr Thr Trp Asp Phe Gly Asp Gly Ser Pro
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Val Leu Thr Gln Ser Gln Pro Ala Ala Asn His Thr Tyr Ala Ser Arg
 1170 1175 1180

Gly Thr Tyr His Val Arg Leu Glu Val Asn Asn Thr Val Ser Gly Ala
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Ala Ala Gln Ala Asp Val Arg Val Phe Glu Glu Leu Arg Gly Leu Ser
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 Val Asp Met Ser Leu Ala Val Glu Gln Gly Ala Pro Val Val Val Ser
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 Ala Ala Val Gln Thr Gly Asp Asn Ile Thr Trp Thr Phe Asp Met Gly
 1235 1240 1245
 Asp Gly Thr Val Leu Ser Gly Pro Glu Ala Thr Val Glu His Val Tyr
 1250 1255 1260
 Leu Arg Ala Gln Asn Cys Thr Val Thr Val Gly Ala Gly Ser Pro Ala
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 Gly His Leu Ala Arg Ser Leu His Val Leu Val Phe Val Leu Glu Val
 1285 1290 1295
 Leu Arg Val Glu Pro Ala Ala Cys Ile Pro Thr Gln Pro Asp Ala Arg
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 Leu Thr Ala Tyr Val Thr Gly Asn Pro Ala His Tyr Leu Phe Asp Trp
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 Thr Phe Gly Asp Gly Ser Ser Asn Thr Thr Val Arg Gly Cys Pro Thr
 1330 1335 1340
 Val Thr His Asn Phe Thr Arg Ser Gly Thr Phe Pro Leu Ala Leu Val
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 Leu Ser Ser Arg Val Asn Arg Ala His Tyr Phe Thr Ser Ile Cys Val
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 Glu Pro Glu Val Gly Asn Val Thr Leu Gln Pro Glu Arg Gln Phe Val
 1380 1385 1390
 Gln Leu Gly Asp Glu Ala Trp Leu Val Ala Cys Ala Trp Pro Pro Phe
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 Pro Tyr Arg Tyr Thr Trp Asp Phe Gly Thr Glu Glu Ala Ala Pro Thr
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 Tyr Leu Val Thr Val Thr Ala Ser Asn Asn Ile Ser Ala Ala Asn Asp
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 Ser Ala Leu Val Glu Val Gln Glu Pro Val Leu Val Thr Ser Ile Lys
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 Val Asn Gly Ser Leu Gly Leu Glu Leu Gln Gln Pro Tyr Leu Phe Ser
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 Gly Gly Trp Leu Glu Gly Pro Glu Val Thr His Ala Tyr Asn Ser Thr
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 Gly Asp Phe Thr Val Arg Val Ala Gly Trp Asn Glu Val Ser Arg Ser
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Val Asn Ala Ser Arg Thr Val Val Pro Leu Asn Gly Ser Val Ser Phe
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 1585 1590 1595 1600
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 1650 1655 1660
 Ser Trp Thr Ala Trp Arg Asp Arg Gly Pro Ala Leu Ala Gly Ser Gly
 1665 1670 1675 1680
 Lys Gly Phe Ser Leu Thr Val Leu Glu Ala Gly Thr Tyr His Val Gln
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 Leu Arg Ala Thr Asn Met Leu Gly Ser Ala Trp Ala Asp Cys Thr Met
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 1745 1750 1755 1760
 Glu Thr Ser Glu Pro Phe Thr Thr His Ser Phe Pro Thr Pro Gly Leu
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 His Leu Val Thr Met Thr Ala Gly Asn Pro Leu Gly Ser Ala Asn Ala
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 Thr Val Glu Val Asp Val Gln Val Pro Val Ser Gly Leu Ser Ile Arg
 1795 1800 1805
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 1810 1815 1820
 Phe Trp Gly Gln Leu Ala Thr Gly Thr Asn Val Ser Trp Cys Trp Ala
 1825 1830 1835 1840
 Val Pro Gly Gly Ser Ser Lys Arg Gly Pro His Val Thr Met Val Phe
 1845 1850 1855
 Pro Asp Ala Gly Thr Phe Ser Ile Arg Leu Asn Ala Ser Asn Ala Val
 1860 1865 1870
 Ser Trp Val Ser Ala Thr Tyr Asn Leu Thr Ala Glu Glu Pro Ile Val
 1875 1880 1885
 Gly Leu Val Leu Trp Ala Ser Ser Lys Val Val Ala Pro Gly Gln Leu
 1890 1895 1900
 Val His Phe Gln Ile Leu Leu Ala Ala Gly Ser Ala Val Thr Phe Arg

1905

1910

1915

1920

Leu Gln Val Gly Gly Ala Asn Pro Glu Val Leu Pro Gly Pro Arg Phe
 1925 1930 1935
 Ser His Ser Phe Pro Arg Val Gly Asp His Val Val Ser Val Arg Gly
 1940 1945 1950
 Lys Asn His Val Ser Trp Ala Gln Ala Val Arg Ile Val Val Leu
 1955 1960 1965
 Glu Ala Val Ser Gly Leu Gln Val Pro Asn Cys Cys Glu Pro Gly Ile
 1970 1975 1980
 Ala Thr Gly Thr Glu Arg Asn Phe Thr Ala Arg Val Gln Arg Gly Ser
 1985 1990 1995 2000
 Arg Val Ala Tyr Ala Trp Tyr Phe Ser Leu Gln Lys Val Gln Gly Asp
 2005 2010 2015
 Ser Leu Val Ile Leu Ser Gly Arg Asp Val Thr Tyr Thr Pro Val Ala
 2020 2025 2030
 Ala Gly Leu Leu Glu Ile Gln Val Arg Ala Phe Asn Ala Leu Gly Ser
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 2050 2055 2060
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 2085 2090 2095
 Gly Asp Gly Ser Pro Gly Gln Asp Thr Asp Glu Pro Arg Ala Glu His
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 Ser Tyr Leu Arg Pro Gly Asp Tyr Arg Val Gln Val Asn Ala Ser Asn
 2115 2120 2125
 Leu Val Ser Phe Phe Val Ala Gln Ala Thr Val Thr Val Gln Val Leu
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 Val Pro Ile Ile Glu Gly Gly Ser Tyr Arg Val Trp Ser Asp Thr Arg
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Asp Leu Val Leu Asp Gly Ser Glu Ser Tyr Asp Pro Asn Leu Glu Asp
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Gly Asp Gln Thr Pro Leu Ser Phe His Trp Ala Cys Val Ala Ser Thr
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Gln Arg Glu Ala Gly Gly Cys Ala Leu Asn Phe Gly Pro Arg Gly Ser
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Ser Thr Val Thr Ile Pro Arg Glu Arg Leu Ala Ala Gly Val Glu Tyr
2325 2330 2335

Thr Phe Ser Leu Thr Val Trp Lys Ala Gly Arg Lys Glu Glu Ala Thr
2340 2345 2350

Asn Gln Thr Val Leu Ile Arg Ser Gly Arg Val Pro Ile Val Ser Leu
2355 2360 2365

Glu Cys Val Ser Cys Lys Ala Gln Ala Val Tyr Glu Val Ser Arg Ser
2370 2375 2380

Ser Tyr Val Tyr Leu Glu Gly Arg Cys Leu Asn Cys Ser Ser Gly Ser
335 2390 2395 2400

Lys Arg Gly Arg Trp Ala Ala Arg Thr Phe Ser Asn Lys Thr Leu Val
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Leu Asp Glu Thr Thr Ser Thr Gly Ser Ala Gly Met Arg Leu Val
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Leu Arg Arg Gly Val Leu Arg Asp Gly Glu Gly Tyr Thr Phe Thr Leu
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Thr Val Leu Gly Arg Ser Gly Glu Glu Gly Cys Ala Ser Ile Arg
2450 2455 2460

Leu Ser Pro Asn Arg Pro Pro Leu Gly Gly Ser Cys Arg Leu Phe Pro
465 2470 2475 2480

Leu Gly Ala Val His Ala Leu Thr Thr Lys Val His Phe Glu Cys Thr
2485 2490 2495

Gly Trp His Asp Ala Glu Asp Ala Gly Ala Pro Leu Val Tyr Ala Leu
2500 2505 2510

Leu Leu Arg Arg Cys Arg Gln Gly His Cys Glu Glu Phe Cys Val Tyr
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Lys Gly Ser Leu Ser Ser Tyr Gly Ala Val Leu Pro Pro Gly Phe Arg
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Pro His Phe Glu Val Gly Leu Ala Val Val Val Gln Asp Gln Leu Gly
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Pro Asn Gly Ser Ala Thr Gly Leu Thr Val Trp Leu His Gly Leu Thr
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Ala Ser Val Leu Pro Gly Leu Leu Arg Gln Ala Asp Pro Gln His Val
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Ile Glu Tyr Ser Leu Ala Leu Val Thr Val Leu Asn Glu Tyr Glu Arg
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 Gln Ile Arg Lys Asn Ile Thr Glu Thr Leu Val Ser Leu Arg Val His
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 Thr Val Asp Asp Ile Gln Gln Ile Ala Ala Ala Leu Ala Gln Cys Met
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 Gly Pro Ser Arg Glu Leu Val Cys Arg Ser Cys Leu Lys Gln Thr Leu
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 His Lys Leu Glu Ala Met Met Leu Ile Leu Gln Ala Glu Thr Thr Ala
 2690 2695 2700
 Gly Thr Val Thr Pro Thr Ala Ile Gly Asp Ser Ile Leu Asn Ile Thr
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 Gly Asp Leu Ile His Leu Ala Ser Ser Asp Val Arg Ala Pro Gln Pro
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 Ser Glu Leu Gly Ala Glu Ser Pro Ser Arg Met Val Ala Ser Gln Ala
 2740 2745 2750
 Tyr Asn Leu Thr Ser Ala Leu Met Arg Ile Leu Met Arg Ser Arg Val
 2755 2760 2765
 Leu Asn Glu Glu Pro Leu Thr Leu Ala Gly Glu Glu Ile Val Ala Gln
 2770 2775 2780
 Gly Lys Arg Ser Asp Pro Arg Ser Leu Leu Cys Tyr Gly Gly Ala Pro
 785 2790 2795 2800
 Gly Pro Gly Cys His Phe Ser Ile Pro Glu Ala Phe Ser Gly Ala Leu
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 Asp Trp Ala Ala Arg Gly His Arg Ser Ser Ala Asn Ser Ala Asn Ser
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2980

2985

2990

Leu Asn Leu Ser Ser His Phe Arg Trp Ser Ala Leu Gln Val Ser Val
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Val Pro Pro Ser His Val Arg Phe Val Phe Pro Glu Pro Thr Ala Asp
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Val Asn Tyr Ile Val Met Leu Thr Cys Ala Val Cys Leu Val Thr Tyr
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Met Val Met Ala Ala Ile Leu His Lys Leu Asp Gln Leu Asp Ala Ser
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Arg Gly Arg Ala Ile Pro Phe Cys Gly Gln Arg Gly Arg Phe Lys Tyr
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Glu Ile Leu Val Lys Thr Gly Trp Gly Arg Gly Ser Gly Thr Thr Ala
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His Val Gly Ile Met Leu Tyr Gly Val Asp Ser Arg Ser Gly His Arg
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His Leu Asp Gly Asp Arg Ala Phe His Arg Asn Ser Leu Asp Ile Phe
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Arg Ile Ala Thr Pro His Ser Leu Gly Ser Val Trp Lys Ile Arg Val
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Trp His Asp Asn Lys Gly Leu Ser Pro Ala Trp Phe Leu Gln His Val
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Ile Val Arg Asp Leu Gln Thr Ala Arg Ser Ala Phe Phe Leu Val Asn
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Asp Trp Leu Ser Val Glu Thr Glu Ala Asn Gly Gly Leu Val Glu Lys
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Glu Val Leu Ala Ala Ser Asp Ala Ala Leu Leu Arg Phe Arg Arg Leu
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Leu Val Ala Glu Leu Gln Arg Gly Phe Phe Asp Lys His Ile Trp Leu
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Ser Arg Leu Ser Pro Leu Ser Val Asp Thr Val Ala Val Gly Leu Val
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Ser Ser Val Val Val Tyr Pro Val Tyr Leu Ala Ile Leu Phe Leu Phe
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Arg Met Ser Arg Ser Lys Val Ala Gly Ser Pro Ser Pro Thr Pro Ala
 345 3350 3355 3360
 Gly Gln Gln Val Leu Asp Ile Asp Ser Cys Leu Asp Ser Ser Val Leu
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 Asp Ser Ser Phe Leu Thr Phe Ser Gly Leu His Ala Glu Gln Ala Phe
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 Val Gly Gln Met Lys Ser Asp Leu Phe Leu Asp Asp Ser Lys Ser Leu
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 Asp Thr His Met Glu Thr Asp Leu Leu Ser Ser Leu Ser Ser Thr Pro
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 Gly Glu Lys Thr Glu Thr Leu Ala Leu Gln Arg Leu Gly Glu Leu Gly
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 Pro Pro Ser Pro Gly Leu Asn Trp Glu Gln Pro Gln Ala Ala Arg Leu
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 Ser Arg Thr Gly Leu Val Glu Gly Leu Arg Lys Arg Leu Leu Pro Ala
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 Trp Cys Ala Ser Leu Ala His Gly Leu Ser Leu Leu Val Ala Val
 3555 3560 3565
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ccaaaaaccgg acggcgtgcg tccgggtggc tttaaaatgg acgacccggac gccgattttgt 6540
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caagaacaca gcgtatccac caatccat aaacgaacgg gcagtcagct tgatctttcc 6840
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ggccggatcc ttcttcttat cgtgctcata ctcctcactg accatagaat ggtcaacgca 6960
ggggccatgc tccgcagcgg cgaccggctg cgggtggatta gcccatcgct cgtccgcagg 7020
gccgttagttc attgaagacg gtcgtgttca acagtagaaaa atttgaatta aagtttttag 7080
aaaagttgaa aatcgagacg tctgtgtgtt aaaaactggaa aaaaatagatg cggaaagagg 7140
cgagctcgcc aatccacgt ctcgttagt cttggagatg cccgattgtt aagagatttc 7200
cgtagatact atgtttatg ggatttcacg tttttgggtt gggacgggtt tttgcata 7260

aacggaaaaa t~~gat~~ggagga ataga~~aa~~acg aa~~at~~gattt gaaactgaaa accatcgast 7320
 atacggcaca at~~cata~~tactac atttatcggg ttatt~~gaa~~ac tgc~~atcc~~aa aagt~~tt~~acaa 7330
 tt~~aa~~att~~ca~~ cataccattt gaagataaca acgaataaaa agacttgaa agg~~gg~~gaaa 7440
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 t~~cg~~agcc~~gc~~ cc~~gagat~~cat aagtttttg t~~ca~~at~~catta~~ atatt~~gat~~tc atctgaat~~ga~~ 7560
 aatt~~gt~~aaaaa tt~~ctt~~taa~~g~~ g~~t~~ttctaaa atatt~~ga~~act gttttttt agattt~~cg~~tt 7620
 agtata~~ta~~at t~~ttt~~gaaaca tacat~~ttt~~tt t~~ttt~~ccaaat t~~tc~~a~~gt~~tatc tt~~t~~acgatt 7680
 t~~tt~~gaaaaat cccaaaaatt gtaa~~ac~~at~~ta~~ aaatt~~ct~~gaa taaac~~gg~~tg~~g~~ aaattt~~gt~~tg~~g~~ 7740
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 aat~~gt~~t~~ct~~ta t~~ttt~~cagagt t~~ca~~acaat~~gc~~ cttagagaaa gttggaaaaat gataat~~gt~~tt 7860
 gtt~~ag~~tat~~at~~ tgagaat~~at~~co at~~g~~ca~~gt~~tg~~a~~ aacaatt~~gt~~ t~~ttt~~ttt~~cg~~ ataacaat~~ta~~ 7920
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 t~~at~~ctt~~c~~cc ct~~tt~~cat~~cc~~ ataat~~gt~~at~~g~~ ccc~~ct~~tt~~ca~~ aat~~gg~~aaaaat at~~gat~~gt~~cg~~g 8040
 ggggagg~~tc~~cc t~~cc~~cc~~tc~~cc cacgacc~~ct~~cc cat 8073

<210> 6
 <211> 815
 <212> PRT
 <213> C. Elegans Pkd-2 protein

<400>	6															
Met	Glu	Gly	Arg	Gly	Glu	Gly	Glu	Asp	L ^e u	P ^r o	P ^r o	T ^h r	S ^e r	T ^y r	P ^h e	
1					5				10					15		
Pro	P ^h e	Glu	Glu	Gly	His	Thr	L ^e u	T ^r p	M ^e t	L ^y s	A ^r g	Glu	L ^y s	I ^l e	L ^y s	
	20						25			30						
His	L ^e u	Gln	Arg	I ^l e	L ^e u	Gln	Phe	His	Ser	A ^s p	Glu	Ser	I ^l e	L ^e u	M ^e t	
	35					40							45			
I ^l e	A ^s p	Lys	Lys	L ^e u	M ^e t	I ^l e	Ser	Gly	Gly	L ^e u	Glu	P ^r o	P ^r o	T ^h r	P ^h e	
	50					55				60						
Cys	V ^a l	L ^e u	A ^s p	A ^r g	Cys	A ^s p	A ⁿ s	H ⁱ s	T ^y r	T ^h r	L ^y s	P ^r o	A ^r g	H ⁱ s		
	65				70				75			80				
L ^e u	P ^r o	P ^r o	Phe	Glu	V ^a l	Phe	L ^e u	Phe	V ^a l	V ^a l	I ^l e	Phe	L ^y s	Cys	Glu	
	85						90					95				
P ^r o	S ^e r	S ^e r	M ^e t	A ⁿ s	T ^y r	Gly	A ^l a	A ^l a	A ^s p	Glu	A ^r g	T ^r p	A ^l a	A ⁿ s	P ^r o	
	100						105					110				
P ^r o	Gln	P ^r o	V ^a l	A ^l a	A ^l a	Glu	H ⁱ s	Gly	P ^r o	S ^e r	Phe	A ^s p	H ⁱ s	S ^e r		
	115				120						125					
M ^e t	V ^a l	S ^e r	Glu	Glu	T ^y r	Glu	H ⁱ s	A ^s p	L ^y s	L ^y s	A ⁿ s	P ^r o	A ^l a	Gln		
	130					135				140						
L ^y s	Glu	Gly	I ^l e	Ser	Phe	Ser	Gln	A ^l a	L ^e u	L ^e u	A ^l a	Ser	Gly	H ⁱ s	Glu	
	145					150					155			160		

Lys Ser Asp Gly Lys Ile Lys Leu Thr Ala Arg Ser Phe Met Glu Val
 155 170 175
 Gly Gly Tyr Ala Val Phe Leu Ile Val Leu Val Tyr Val Ala Phe Ala
 180 185 190
 Gln Asn Ser Ile Gln Ser Tyr Tyr Ser Lys Val Met Ser Asp Leu
 195 200 205
 Phe Val Ala Ser Thr Gly Ala Ser Gly Ala Pro Ala Phe Gly Ser Cys
 210 215 220
 Thr Ser Met Asp Asn Ile Trp Asp Trp Leu Ser Gln Val Leu Ile Pro
 225 230 235 240
 Gly Ile Tyr Trp Thr Glu Thr Ser Asn Ser Thr Asp Asn Glu Asn Met
 245 250 255
 Ile Tyr Tyr Glu Asn Arg Leu Leu Gly Glu Pro Arg Ile Arg Met Leu
 260 265 270
 Lys Val Thr Asn Asp Ser Cys Thr Val Met Lys Ser Phe Gln Arg Glu
 275 280 285
 Ile Lys Glu Cys Phe Ala Asn Tyr Glu Glu Lys Leu Glu Asp Lys Thr
 290 295 300
 Met Val Gly Asp Gly Ser Val Asp Ala Phe Ile Tyr Ala Thr Ala Lys
 305 310 315 320
 Glu Leu Glu Asn Leu Lys Thr Val Gly Thr Ile Ala Ser Tyr Gly
 325 330 335
 Gly Gly Phe Val Gin Arg Leu Pro Val Ala Gly Ser Thr Glu Ala Gln
 340 345 350
 Ser Ala Ile Ala Thr Leu Lys Ala Asn Arg Trp Ile Asp Arg Gly Ser
 355 360 365
 Arg Ala Ile Ile Val Asp Phe Ala Leu Tyr Asn Ala Asn Ile Asn Leu
 370 375 380
 Phe Cys Val Val Lys Leu Leu Phe Glu Leu Pro Ala Ser Gly Gly Val
 385 390 395 400
 Ile Thr Thr Pro Lys Leu Met Thr Tyr Asp Leu Leu Thr Tyr Gln Thr
 405 410 415
 Ser Gly Gly Thr Arg Met Met Ile Phe Glu Gly Ile Phe Cys Gly Phe
 420 425 430
 Ile Leu Tyr Phe Ile Phe Glu Leu Phe Ala Ile Gly Arg His Arg
 435 440 445
 Leu His Tyr Leu Thr Gln Phe Trp Asn Leu Val Asp Val Val Leu Leu
 450 455 460
 Gly Phe Ser Val Ala Thr Ile Ile Leu Ser Val Asn Arg Thr Lys Thr
 465 470 475 480
 Gly Val Asn Arg Val Asn Ser Val Ile Glu Asn Gly Leu Thr Asn Ala
 485 490 495
 Pro Phe Asp Asp Val Thr Ser Ser Glu Asn Ser Tyr Leu Asn Ile Lys
 500 505 510
 Ala Cys Val Val Phe Val Ala Trp Val Lys Val Phe Lys Phe Ile Ser

515

520

525

Val Asn Lys Thr Met Ser Gln Leu Ser Ser Thr Leu Thr Arg Ser Ala
 530 535 540

Lys Asp Ile Gly Gly Phe Ala Val Met Phe Ala Val Phe Phe Phe Ala
 545 550 555 560

Phe Ala Gln Phe Gly Tyr Leu Cys Phe Gly Thr Gln Ile Ala Asp Tyr
 565 570 575

Ser Asn Leu Tyr Asn Ser Ala Phe Ala Leu Leu Arg Leu Ile Leu Gly
 580 585 590

Asp Phe Asn Phe Ser Ala Leu Glu Ser Cys Asn Arg Phe Phe Gly Pro
 595 600 605

Ala Phe Phe Ile Ala Tyr Val Phe Phe Val Ser Phe Ile Leu Leu Asn
 610 615 620

Met Phe Leu Ala Ile Ile Asn Asp Ser Tyr Val Glu Val Lys Ala Glu
 625 630 635 640

Leu Ala Arg Lys Lys Asp Gly Glu Gly Ile Leu Asp Trp Phe Met Asn
 645 650 655

Lys Val Arg Gly Leu Thr Lys Arg Gly Lys Arg Pro Asp Ala Pro Gly
 660 665 670

Glu Asp Ala Thr Tyr Glu Asp Tyr Lys Leu Met Leu Tyr Arg Ala Gly
 675 680 685

Tyr Ala Glu Lys Asp Ile Asn Glu Ala Phe Thr Arg Phe Asn Val Thr
 690 695 700

Ser Met Thr Glu His Val Pro Glu Lys Val Ala Glu Asp Ile Ala Asp
 705 710 715 720

Glu Val Ala Arg Met Thr Glu Gln Lys Arg Asn Tyr Met Glu Asn His
 725 730 735

Arg Asp Tyr Ala Asn Leu Asn Arg Arg Val Asp Gln Met Gln Glu Ser
 740 745 750

Val Phe Ser Ile Val Asp Arg Ile Glu Gly Val Asn Ala Thr Leu Gln
 755 760 765

Thr Ile Glu Lys Gln Arg Val Gln Gln Gln Asp Gly Gly Asn Leu Met
 770 775 780

Asp Leu Ser Ala Leu Leu Thr Asn Gln Val Arg Asn Arg Glu Ser Ala
 785 790 795 800

Ala Arg Arg Pro Thr Ile Thr Ser Ile Ala Asp Lys Lys Glu Glu
 805 810 815

<210> 7

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Outside primer for PCR screening of
 lov-1 genomic (sy582) deletion

<400> 7

ctctatttgt ggttcggtgg cg

22

<210> 8
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Outside primer for PCR screening of
lov 1 genomic (sy582) deletion

<400> 8
gggtttttcc gttttcatgg gg

22

<210> 9
<211> 22
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Nested primer for PCR screening of
lov 1 genomic (sy582) deletion

<400> 9
cttggaccgga tgcaaacagcg ag

22

<210> 10
<211> 22
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Nested primer for PCR screening of
lov-1 genomic (sy582) deletion

<400> 10
aacgttgatt ggttcaagtg tg

22

<210> 11
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Outside primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 11
ccccctcggtt gaccattcta tgg

23

<210> 12
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Outside primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 12

acgtgatcc ctgtcgatcc ag

22

<210> 13

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Nested primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 13

agatcaagct gactgcgggt tc

22

<210> 14

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Nested primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 14

gatccagcga ttagccttta acg

23

<210> 15

<211> 2870

<212> PRT

<213> C. Elegans Lov-1 sy582 deletion protein

<400> 15

Met Val Leu Arg Phe Ser Pro Pro Phe Arg Phe Ser Thr Thr Ser Phe
1 5 10 15

Phe Ser Cys Cys Leu Phe Cys Ser Glu Phe Ile Phe Val Phe Arg Arg
20 25 30

Ile Phe Thr Lys Leu Leu Gln Asp Asn Leu Pro Ala His Trp Met Lys
35 40 45

Lys Ser Asn Phe Phe Val Leu Leu Leu Ala Ile Ser Ala Ile Gln
50 55 60

Ile Asp Gly Leu His Tyr Gln Leu Leu Asp Gly Ile Ala Thr Phe Arg
65 70 75 80

Leu Asp Asn Asp Asp Thr Thr Ile Gly Gly Val Pro Arg Asn Ser Gln
85 90 95

Gly Val Val Lys Ile Lys Leu Ser Cys Gly Leu Asn Arg Leu Ser Val
100 105 110

Glu Asn Lys Val Thr Glu Val Ser Ser Leu Glu Leu Ile His Asn Cys
115 120 125

Ile Gln Thr Glu Thr Arg Leu Val Gly Leu Phe Leu Asn Ser Thr Trp
130 135 140

Ile Thr Leu Asn Glu Val Asn Asp Asp Glu Ile Ser Ile Ala Val
145 150 155 160

Glu Ala Lys Tyr Glu Val Cys Tyr Asp Asp Gly Ile Asp Arg Cys Asp

165

170

175

Gly Ser Leu Trp Trp Leu Gln Val Gly Gly Asn Glu Met Ala Leu Leu
 180 185 190

Gly Tyr Arg Glu Lys Cys Glu Ser Gly Glu Ile Asn Glu Glu Tyr Ala
 195 200 205

Arg Arg Met Cys Lys Arg Pro Tyr Arg Ser Glu Lys Ser Thr Ala Ile
 210 215 220

Ser Asp Ser Gln Gly Val Tyr Tyr Asp Gly Gln Val Leu Lys Gly Val
 225 230 235 240

Arg Ala Lys Gln Phe Ser Met Arg Thr Ser Gly Ser Pro Thr Leu Arg
 245 250 255

Arg Met Lys Arg Asp Ala Gly Asp Asn Thr Cys Asp Tyr Thr Ile Glu
 260 265 270

Ser Thr Ser Thr Ser Thr Thr Pro Thr Thr Thr Thr Val Thr Ser
 275 280 285

Thr Val Thr Ser Thr Thr Val Pro Thr Ser Thr Ser Thr Val Thr
 290 295 300

Thr Ala Met Ser Thr Ser Thr Ser Pro Ser Thr Ser Thr Thr Ile
 305 310 315 320

Glu Ser Thr Ser Thr Thr Phe Thr Ser Thr Ala Ser Thr Ser Thr Ser
 325 330 335

Ser Thr Ser Thr Thr Gln Gln Ser Ser Ser Thr Ile Thr Ser Ser Pro
 340 345 350

Ser Ser Thr Thr Leu Ser Thr Ser Ile Pro Thr Thr Thr Thr Pro Glu
 355 360 365

Ile Thr Ser Thr Leu Ser Ser Leu Pro Asp Asn Ala Ile Cys Ser Tyr
 370 375 380

Leu Asp Glu Thr Thr Thr Ser Thr Thr Phe Thr Thr Thr Met Leu Thr
 385 390 395 400

Ser Thr Thr Thr Glu Glu Pro Ser Thr Ser Thr Thr Thr Thr Glu Val
 405 410 415

Thr Ser Thr Ser Ser Thr Val Thr Thr Glu Pro Thr Thr Thr Leu
 420 425 430

Thr Thr Ser Thr Ala Ser Thr Ser Thr Glu Pro Ser Thr Ser Thr
 435 440 445

Val Thr Thr Ser Pro Ser Thr Ser Pro Val Thr Ser Thr Val Thr Ser
 450 455 460

Ser Ser Ser Ser Ser Thr Thr Val Thr Thr Pro Thr Ser Thr Glu Ser
 465 470 475 480

Thr Ser Thr Ser Pro Ser Ser Thr Val Thr Thr Ser Thr Thr Ala Pro
 485 490 495

Ser Thr Ser Thr Thr Gly Pro Ser Ser Ser Ser Ser Thr Pro Ser Ser
 500 505 510

Thr Ala Ser Ser Ser Val Ser Ser Thr Ala Ser Ser Thr Gln Ser Ser

515

520

525

Thr Ser Thr Gln Gln Ser Ser Thr Thr Lys Ser Glu Thr Thr Thr
 530 535 540
 Ser Ser Asp Gly Thr Asn Pro Asp Phe Tyr Phe Val Glu Lys Ala Thr
 545 550 555 560
 Thr Thr Phe Tyr Asp Ser Thr Ser Val Asn Leu Thr Leu Asn Ser Gly
 565 570 575
 Leu Gly Ile Ile Gly Tyr Gln Thr Ser Ile Glu Cys Thr Ser Pro Thr
 580 585 590
 Ser Ser Asn Tyr Val Ser Thr Thr Lys Asp Gly Ala Cys Phe Thr Lys
 595 600 605
 Ser Val Ser Met Pro Arg Leu Gly Gly Thr Tyr Pro Ala Ser Thr Phe
 610 615 620
 Val Gly Pro Gly Asn Tyr Thr Phe Arg Ala Thr Met Thr Thr Asp Asp
 625 630 635 640
 Lys Lys Val Tyr Tyr Thr Tyr Ala Asn Val Tyr Ile Gln Glu Tyr Ser
 645 650 655
 Ser Thr Thr Ile Glu Ser Glu Ser Ser Thr Ser Ala Val Ala Ser Ser
 660 665 670
 Thr Ser Ser Thr Pro Ser Thr Pro Ser Ser Thr Leu Ser Thr Ser Thr
 675 680 685
 Val Thr Glu Pro Ser Ser Thr Arg Ser Ser Asp Ser Thr Thr Thr Ser
 690 695 700
 Ala Gly Ser Thr Thr Leu Gln Glu Ser Thr Thr Thr Ser Glu Glu
 705 710 715 720
 Ser Thr Thr Asp Ser Ser Thr Thr Ile Ser Asp Thr Ser Thr Ser
 725 730 735
 Thr Ser Ser Pro Ser Ser Thr Ala Asp Ser Thr Ser Thr Leu Ser
 740 745 750
 Val Asp Gln Phe Asp Phe Ile Leu Asp Ser Gly Leu Ser Trp Asn Glu
 755 760 765
 Thr Arg His Asn Glu Asp Ser Ile Asn Ile Val Pro Leu Pro Thr Asn
 770 775 780
 Ala Ile Thr Pro Thr Glu Arg Ser Gln Thr Phe Glu Cys Arg Asn Val
 785 790 795 800
 Ser Thr Glu Pro Phe Leu Ile Ile Lys Glu Ser Thr Cys Leu Asn Tyr
 805 810 815
 Ser Asn Thr Val Leu Asn Ala Thr Tyr Ser Ser Asn Ile Pro Ile Gln
 820 825 830
 Pro Ile Glu Thr Phe Leu Val Gly Ile Gly Thr Tyr Glu Phe Arg Ile
 835 840 845
 Asn Met Thr Asp Leu Thr Thr Met Gln Val Val Ser His Ile Phe Thr
 850 855 860
 Leu Asn Val Val Ala Asp Ser Thr Ser Thr Ser Glu Val Thr Ser Thr

865	870	875	880
Thr Ser Thr Gly Ser Ser Ser Glu Ser Ser Ala Ile Ser Thr Thr Ser			
885	890	895	900
Gly Ile Glu Ser Thr Ser Thr Leu Glu Ala Ser Thr Thr Asp Ala Ser			
900	905	910	915
Gln Asp Ser Ser Thr Ser Thr Asp Ser Gly Thr Thr Ser Asp Ser			
915	920	925	930
Thr Thr Ile Asp Ser Ser Asn Ser Thr Pro Ser Thr Ser Asp Ser Ser			
930	935	940	945
Gly Leu Ser Gln Thr Pro Ser Asp Ser Ser Ser Ala Ser Asp Ser Met			
945	950	955	960
Arg Thr Thr Thr Val Asp Pro Asp Ala Ser Thr Glu Thr Pro Tyr Asp			
965	970	975	980
Phe Val Leu Glu Asn Leu Thr Trp Asn Glu Thr Val Tyr Tyr Ser Glu			
980	985	990	995
Asn Pro Phe Tyr Ile Thr Pro Ile Pro Asn Lys Glu Pro Gly Ala Leu			
995	1000	1005	1010
Thr Thr Ala Met Thr Cys Gln Cys Arg Asn Asp Ser Ser Gln Pro Phe			
1010	1015	1020	1025
Val Leu Leu Lys Glu Ser Asn Cys Leu Thr Glu Phe Gly Lys Asn Gly			
1025	1030	1035	1040
Ala Tyr Ser Ala Ser Val Ser Phe Asn Pro Met Thr Ser Phe Val Pro			
1045	1050	1055	1060
Ala Thr Gly Thr Tyr Glu Phe Leu Ile Asn Val Thr Asn Arg Ala Ser			
1060	1065	1070	1075
Gly Glu Ser Ala Ser His Ile Phe Thr Met Asn Val Val Leu Pro Thr			
1075	1080	1085	1090
Thr Thr Thr Glu Thr Pro Pro Thr Thr Val Ser Ser Ser Asp Asp Ala			
1090	1095	1100	1105
Gly Gly Lys Thr Gly Gly Thr Gly Ala Thr Gly Gly Thr Gly Gly Thr			
1110	1115	1120	1125
Gly Ser Gly Gly Ser Ala Thr Thr Leu Ser Thr Gly Asp Ala Val Arg			
1125	1130	1135	1140
Ser Thr Thr Ser Gly Ser Gly Gln Ser Ser Thr Gly Ser Gly			
1140	1145	1150	1155
Ala Gly Gly Ser Gly Thr Thr Ala Ser Gly Ser Gly Ser Gly Ser			
1155	1160	1165	1170
Ser Gly Thr Gly Ser Asp Gly Val Asn Ser Gly Lys Thr Thr Ala Leu			
1170	1175	1180	1185
Asn Gly Asp Gly Thr Gly Ser Gly Thr Ala Thr Thr Pro Gly Ser His			
1190	1195	1200	1205
Leu Gly Asp Gly Gly Ser Thr Ser Gly Ser Gly Ser Asp Ser Asn Gly			
1205	1210	1215	1220
Ser Ser Gly Val Ser Thr Lys Ser Ser Ser Gly Ser Asp Thr Ser Gly			

1220 1225 1230

Ser Ser Asp Ser Ser Gly Ala Asn Gly Ala Phe Ser Ala Thr Ala Gln
 1235 1240 1245

Pro Ser Thr Arg Thr Thr Lys Thr Arg Ser Ser Leu Ala Thr Val Ser
 1250 1255 1260

Pro Ile Ser Ala Ala Glu Gln Ala Ile Ile Asp Ala Gln Lys Ala Asp
 1265 1270 1275 1280

Val Met Asn Gln Leu Ala Gly Ile Met Asp Gly Ser Ala Ser Asn Asn
 1285 1290 1295

Ser Leu Asn Thr Ser Ser Leu Leu Asn Gln Ile Ser Ser Leu Pro
 1300 1305 1310

Ala Ala Asp Leu Val Glu Val Ala Gln Ser Leu Leu Ser Asn Thr Leu
 1315 1320 1325

Lys Ile Pro Gly Val Gly Asn Met Ser Ser Val Asp Val Leu Lys Thr
 1330 1335 1340

Leu Gln Asp Asn Ile Ala Thr Thr Asn Ser Glu Leu Ala Asp Glu Met
 1345 1350 1355 1360

Ala Lys Val Ile Thr Lys Leu Ala Asn Val Asn Met Thr Ser Ala Gln
 1365 1370 1375

Ser Leu Asn Ser Val Leu Ser Ser Leu Asp Leu Ala Leu Lys Gly Ser
 1380 1385 1390

Thr Val Tyr Thr Leu Gly Val Ser Ser Thr Lys Ser Lys Asp Gly Thr
 1395 1400 1405

Tyr Ala Val Ile Phe Gly Tyr Val Ile Ala Ser Gly Tyr Thr Leu Val
 1410 1415 1420

Ser Pro Arg Cys Thr Leu Ser Ile Tyr Gly Ser Thr Ile Tyr Leu Thr
 1425 1430 1435 1440

Gly Asp Thr Arg Ala Ser Tyr Lys Gln Leu Asp Gly Asp Thr Val Thr
 1445 1450 1455

Ala Asp Thr Met Leu Ala Ala Ile Gly Ile Gln Gly Met Phe Ala
 1460 1465 1470

Thr Asn Gly Arg Thr Val Gln Val Glu Gln Asp Lys Ile Asp Asp Lys
 1475 1480 1485

Arg Ser Leu Val Ser Gly Asn Ile Met Ala Thr Met Ser Gly Val Gly
 1490 1495 1500

Asp Val Gln Ser Gly Glu Tyr Ser Tyr Asn Asp Met Tyr Val Thr Ala
 1505 1510 1515 1520

Trp Asn Val Thr Tyr Asp Asn Ser Thr Val Gly Ser Thr Ser Gln Lys
 1525 1530 1535

Asn Thr Ser Phe Ser Phe Asn Ile Pro Val Ser Glu Val Gln Tyr Ile
 1540 1545 1550

Leu Leu Ile Glu Ser Gly Thr Met Ile Lys Leu His Ser Thr Gln Asn
 1555 1560 1565

Ile Val Ser Arg Gly Leu Val Val Thr Ala Ser Tyr Gly Gly Val Thr
 1570 1575 1580

Tyr Thr Ile Thr Cys Thr Asn Gly Thr Gly Lys Phe Val Glu Val Asp
 1535 1590 1595 1600
 Thr Asp Asn Ala Ile Phe Ser Tyr Asn Ala Asp Ser Phe Thr Val Val
 1605 1610 1615
 Ala Ser Asp Gly Ser Ser Ala Ser Thr Val Lys Lys Leu Ile Gln Met
 1620 1625 1630
 Pro Ile Val Ile Glu Asn Val Asn Leu Ala Leu Phe Asn Gln Thr Thr
 1635 1640 1645
 Ser Pro Leu Val Phe Ser Asn Ala Gly Ser Tyr Ser Met Arg Met Val
 1650 1655 1660
 Leu Ser Pro Gln Asp Ile Gly Ile Pro Ala Val Ser Ala Leu Ser Gln.
 1665 1670 1675 1680
 Thr Val Ser Ile Ser Thr Leu Ser Pro Thr Ala Ser Tyr Thr Lys Asp
 1685 1690 1695
 Asp Leu Gln Ser Leu Ile Lys Glu Gln Thr Leu Val Thr Val Ser Gly
 1700 1705 1710
 Thr Thr Ser Asn Ser Leu Leu Ser Ile Ala Gly Ser Leu Thr Ser Ala
 1715 1720 1725
 Leu Lys Ile Ala Leu Asp Asn Pro Leu Ser Ser Asp Leu Ala Ala Asn
 1730 1735 1740
 Leu Lys Tyr Ala Thr Asp Asn Tyr Asp Ser Leu Tyr Asn Val Leu Pro
 1745 1750 1755 1760
 Ser Asp Pro Asp Asn Ile Val Tyr Val Glu Glu Met Thr Ser Glu Glu
 1765 1770 1775
 Trp Ala Ala Tyr Val Thr Lys Met Phe Gln Lys Asn Ile Ala Lys Asn
 1780 1785 1790
 Leu Ala Asn Gln Leu Ala Ser Thr Leu Asp Thr Leu Glu Asn Thr Leu
 1795 1800 1805
 Ala Ala Arg Ala Ile Ala Thr Gly Asn Leu Pro Tyr Asp Tyr Ser Asn
 1810 1815 1820
 Ser Val Asp Gly Thr Gly Met Val Ile Val Ile Asp Asp Ala Ser Asn
 1825 1830 1835 1840
 Ile Val Gly Lys Thr Gln Asn Cys Glu Glu Trp Ala Phe Lys Leu Pro
 1845 1850 1855
 Ser Pro Ala Ser Thr Leu Asn Thr Ala Glu Ile Thr Asp Lys Thr Leu
 1860 1865 1870
 Ile Gln Val Gly Leu Val Cys Tyr Ala Thr Asn Pro Arg Thr Tyr Val
 1875 1880 1885
 Asp Asn Phe Asp Met Leu Ile Thr Ser Gly Ala Leu Glu Ala His Ile
 1890 1895 1900
 Lys Asp Glu Asn Gln Ile Ile Ile Pro Ile Thr Gly Thr Thr Ala Pro
 1905 1910 1915 1920
 Ile Tyr Val Asn Gly Arg Gly Ser Glu Asp Asp Ala Val Leu Thr Leu
 1925 1930 1935

Met Gln Gln Gly Asp Phe Ala Ser Tyr Gln Ile Leu Asp Leu His Ala
 1943 1945 1950
 Phe Arg Thr Thr Asn Trp Asn Asn Ser Leu Gln Val Glu Ile Ile Ala
 1955 1950 1965
 Ser Gln Asp Tyr Glu Ile Pro Asn Asn Asp Asp Thr Tyr Met Phe Ser
 1970 1975 1980
 Ser Phe Gln Ser Leu Pro Gly Pro Leu Glu Ser Asn His Glu Trp Ile
 1985 1990 1995 2000
 Phe Asp Leu Asn Thr Leu Asn Lys Thr Ser Asn Tyr Phe Val Thr Ala
 2005 2010 2015
 Gly Asn Leu Ile Asn Asn Thr Gly Leu Phe Phe Ile Gly Ile Gly Lys
 2020 2025 2030
 Arg Asn Ser Ser Thr Asn Thr Gly Asn Ser Ser Asp Ile Val Asn Tyr
 2035 2040 2045
 Gly Gln Tyr Asp Ser Met Gln Trp Ser Phe Ala Arg Ser Val Pro Met
 2050 2055 2060
 Asp Tyr Gln Val Ala Ala Val Ser Lys Gly Cys Tyr Phe Tyr Gln Lys
 2065 2070 2075 2080
 Thr Ser Asp Val Phe Asn Ser Gln Gly Met Tyr Pro Ser Asp Gly Gln
 2085 2090 2095
 Gly Met Gln Phe Val Asn Cys Ser Thr Asp His Leu Thr Met Phe Ser
 2100 2105 2110
 Val Gly Ala Phe Asn Pro Thr Ile Asp Ala Asp Phe Ser Tyr Asn Tyr
 2115 2120 2125
 Asn Val Asn Glu Ile Glu Lys Asn Val Lys Val Met Ile Ala Ala Val
 2130 2135 2140
 Phe Met Leu Val Val Tyr Gly Cys Leu Thr Ile Asn Ala Ile Ile Cys
 2145 2150 2155 2160
 Gln Arg Lys Asp Ala Ser Arg Gly Arg Leu Arg Phe Leu Lys Asp Asn
 2165 2170 2175
 Glu Pro His Asp Gly Tyr Met Tyr Val Ile Ala Val Glu Thr Gly Tyr
 2180 2185 2190
 Arg Met Phe Ala Thr Thr Asp Ser Thr Ile Cys Phe Asn Leu Ser Gly
 2195 2200 2205
 Asn Glu Gly Asp Gln Ile Phe Arg Ser Phe Arg Ser Glu Glu Asp Gly
 2210 2215 2220
 Asn Trp Glu Phe Pro Phe Ser Trp Gly Thr Thr Asp Arg Phe Val Met
 2225 2230 2235 2240
 Thr Thr Ala Phe Pro Leu Gly Glu Leu Glu Tyr Met Arg Leu Trp Leu
 2245 2250 2255
 Asp Asp Ala Gly Leu Asp His Arg Glu Ser Trp Tyr Cys Asn Arg Ile
 2260 2265 2270
 Ile Val Lys Asp Leu Gln Thr Gln Asp Ile Tyr Tyr Phe Pro Phe Asn
 2275 2280 2285
 Asn Trp Leu Gly Thr Lys Asn Gly Asp Gly Glu Thr Glu Arg Leu Ala

2290	2295	2300
Arg Val Glu Tyr Lys Arg Arg Phe Leu Asp Glu Ser Met Ser Met His		
2305	2310	2315
Met Leu Ala Gln Thr Ile Ser Trp Phe Ala Met Phe Thr Gly Gly Gly		
2325	2330	2335
Asn Arg Leu Arg Asp Arg Val Ser Arg Gln Asp Tyr Ser Val Ser Ile		
2340	2345	2350
Ile Phe Ser Leu Val Val Ser Met Ile Ser Ile Thr Ile Leu Lys		
2355	2360	2365
Ser Asp Asn Ser Ile Ile Ser Asp Ser Lys Ser Val Ser Glu Phe Thr		
2370	2375	2380
Phe Thr Ile Lys Asp Ile Ala Phe Gly Val Gly Phe Gly Val Leu Ile		
2385	2390	2395
Thr Phe Leu Asn Ser Leu His Ile Leu Leu Cys Thr Lys Cys Arg Ser		
2405	2410	2415
His Ser Glu His Tyr Tyr Lys Lys Arg Lys Arg Glu Asp Pro Glu		
2420	2425	2430
Phe Lys Asp Asn Ser Gly Ser Trp Pro Met Phe Met Ala Gly Met Ala		
2435	2440	2445
Arg Thr Ile Ile Val Phe Pro Val Leu Met Gly Leu Ile Tyr Ile Ser		
2450	2455	2460
Gly Ala Gly Met Ser Leu Met Asp Asp Leu Ala Asn Ser Phe Tyr Ile		
2465	2470	2475
Arg Phe Leu Ile Ser Leu Ile Leu Trp Ala Val Val Phe Glu Pro Ile		
2485	2490	2495
Lys Gly Leu Ile Trp Ala Phe Leu Ile Leu Lys Thr Arg Lys Ser His		
2500	2505	2510
Lys Ile Ile Asn Lys Leu Glu Gly Ser Asp Gly Thr Val Val Lys Tyr		
2515	2520	2525
Tyr Glu Met Leu Tyr Ile Phe Phe Ser Val Leu Ile Phe Val Lys Glu		
2530	2535	2540
Ile Val Phe Tyr Leu Tyr Gly Arg Tyr Lys Val Ile Thr Thr Met Lys		
2545	2550	2555
Pro Thr Arg Asn Pro Phe Lys Ile Val Tyr Gln Leu Ala Leu Gly Asn		
2565	2570	2575
Phe Ser Pro Trp Asn Phe Met Asp Leu Ile Val Gly Ala Leu Ala Val		
2580	2585	2590
Ala Ser Val Leu Ala Tyr Thr Ile Arg Gln Arg Thr Thr Asn Arg Ala		
2595	2600	2605
Met Glu Asp Phe Asn Ala Asn Asn Gly Asn Ser Tyr Ile Asn Leu Thr		
2610	2615	2620
Glu Gln Arg Asn Trp Glu Ile Val Phe Ser Tyr Cys Leu Ala Gly Ala		
2625	2630	2635
Val Phe Phe Thr Ser Cys Lys Met Ile Arg Ile Leu Arg Phe Asn Arg		

2645

2650

2655

Arg Ile Gly Val Leu Ala Ala Thr Leu Asp Asn Ala Leu Gly Ala Ile
 2660 2665 2670

Val Ser Phe Gly Ile Ala Phe Leu Phe Phe Ser Met Thr Phe Asn Ser
 2675 2680 2685

Val Leu Tyr Ala Val Leu Gly Asn Lys Met Gly Gly Tyr Arg Ser Leu
 2690 2695 2700

Met Ala Thr Phe Gln Thr Ala Leu Ala Gly Met Leu Gly Lys Leu Asp
 2705 2710 2715 2720

Val Thr Ser Ile Gln Pro Ile Ser Gln Phe Ala Phe Val Val Ile Met
 2725 2730 2735

Ile Tyr Met Ile Ala Gly Ser Lys Leu Val Leu Gln Leu Tyr Val Thr
 2740 2745 2750

Ile Ile Met Phe Glu Phe Glu Glu Ile Arg Asn Asp Ser Glu Lys Gln
 2755 2760 2765

Thr Asn Asp Tyr Glu Ile Ile Asp His Ile Lys Tyr Lys Thr Lys Arg
 2770 2775 2780

Arg Leu Gly Leu Leu Glu Pro Lys Asp Phe Ala Pro Val Ser Ile Ala
 2785 2790 2795 2800

Asp Thr Gln Lys Asp Phe Arg Leu Phe His Ser Ala Val Ala Lys Val
 2805 2810 2815

Asn Leu Leu His His Arg Ala Thr Arg Met Leu Gln Thr Gln Gly Gln
 2820 2825 2830

Tyr Gln Asn Gln Thr Val Ile Asn Tyr Thr Leu Ser Tyr Asp Pro Val
 2835 2840 2845

Ser Ala Ile His Glu Thr Gly Pro Lys Arg Phe Gln Lys Trp Arg Leu
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Asn Asp Val Glu Lys Asp
 2865 2870

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 <211> 200
 <212> PRT
 <213> C. Elegans Pkd-2 deletion mutant (sy606) protein

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Pro Phe Glu Glu Gly His Thr Leu Trp Met Lys Arg Glu Lys Ile Lys
 20 25 30

His Leu Gln Arg Ile Leu Gln Phe His Ser Asp Glu Ser Ile Leu Met
 35 40 45

Ile Asp Lys Lys Leu Met Ile Ser Gly Gly Leu Glu Pro Pro Thr Phe
 50 55 60

Cys Val Leu Asp Arg Cys Asp Asn His Tyr Thr Thr Lys Pro Arg His
 65 70 75 80

Leu Pro Pro Phe Glu Val Phe Leu Phe Val Val Ile Phe Lys Cys Glu
85 90 95

Pro Ser Ser Met Asn Tyr Gly Ala Ala Asp Glu Arg Trp Ala Asn Pro
100 105 110

Pro Gln Pro Val Ala Ala Ala Glu His Gly Pro Ser Phe Asp His Ser
115 120 125

Met Val Ser Glu Glu Tyr Glu His Asp Lys Lys Asn Pro Ala Gln
130 135 140

Lys Glu Gly Ile Ser Phe Ser Gln Ala Leu Leu Ala Ser Gly His Glu
145 150 155 160

Lys Ser Asp Gly Lys Ile Lys Leu Thr Ala Arg Ser Phe Met Glu Val
165 170 175

Gly Gly Tyr Ala Val Phe Leu Ile Val Leu Val Tyr Asp Ser Ser Thr
180 185 190

Pro Arg Gln Lys Ser Leu Lys Thr
195 200